SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO:	Nucleotide location of first	ł .	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19160	49528	A	19269	351	509	VVPLPLPPLDSWTCESWLRSHI LTGESENPDHKRI*PYTSDLPEG
10161	10.550	<b>↓</b>	10070	701	4224	PGPWATD
19161	49529	Α	19270	781	4324	ESVREKVLCGLIQQTVTSVC/W
						SGTYIFAVLLVCVVFHSGAQEK NYTIREEIPENVLIGNLLKDLNL
					:	SLIPNKSLTTTMQFKLVYKTGD
						VPLIRIEEDTGEIFTTGARIDREK
				]		LCAGIPRDEHCFYEVEVAILPDE
						IFRLVKIRFLIEDINDNAPLFPAT
	:			1		VINISIPENSAINSKYTLPAAVDP
	ļ					DVGINGVQNYELIKSQNIFGLD
						VIETPGGDKMPQLIVQKELDRE
						EKDTYVMKVKVEDGGFPQRSS
19162	49530	A	19271	3	644	FRSLPRRCLPRWQPSSFLGHHG
1,710-	.,,					PPFRPPCEAPKIVKKENPRKFIR
						APSQTGY\AKLSVNW\RKTPEAI
						DN\RVR\RRF\KGQILMP\NIGYG
						SNKK\TKHM\LPSGFRKFLV\HN
						VKEPEIAADVATNLTVPRSLTM
						FPPRNRKAIRGKGACPIGPFRSS
						PNPIARVAQLKEN*VRQLMCN
						KSYCAEIAHNVSSKNRKAIVER
						AAQLAIRVTNPNARAQ
19163	49531	В	19272	27	161	
19164	49532	Α	19273	80	430	RWLATHYLTVKLSLLPNLAVT
						CQPKREQNITSNESTDC*VIVTS TKSDSLYTVGMLALSVRAIRCP
						LYLLTGLIFVSKNDLWYCELQS
						HYTVVLLLFMRKRSRFQSNVQ
						RKLMTNF
19165	49533	A	19274	482	832	RWLATHYLTVKLSLLPNLAVT
15105	1,7333		''	1.02	032	CQPKREQNITSNESTDC*VIVTS
						TKSDSLYTVGMLALSVRAIRCP
				:		LYLLTGLIFVSKNDLWYCELQS
		1				HYTVVLLLFMRKRSRFQSNVQ
						RKLMTNF
19166	49534	A	19275	121	201	SMYMDEN*RLELITL*FSSMYM
		l				DHSLL
19167	49535	В	19276	1	1147	
19168	49536	Α	19277	305	442	AGLFKSRKVSSICIRFE*ILTSCL
						GVRISTGKLVARIHGSFADAE
19169	49537	Α	19278	133	527	GFLERPRILIKVTLSIKQKSTSIF
						VGGQFSKFRFLTVLAWKCLSQP
						CF*A*TPASQHVS*SIRAISSMIS
						KRSELLVRLWRRPSP*AGLFKS
						RKVSSICIRFE*ILTSCLGVRIST
10150	10520	<del>                                     </del>	10020	521	660	GKLVARTHGSFADAE
19170	49538	A	19279	531	669	AGLFKSRKVSSICIRFE*ILTSCL
		ل	L	<u> </u>	L	GVRISTGKLVARTHGSFADAE

SEQ ID NO:	SEQ ID NO: of peptide	Met	SEQ ID NO:	Nucleotide location of first		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
NO:	sequence	nou	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19171	49539	Α	19280	80	428	RWLATPYLTVKLSLLPNLAVTC
				•		QPKREQNITSNESTDC*VIVTST
						KSDSLYTVGMLALSVRAIRCPL
						YLLTGLISVSKNGLWYCELQSH
1					1	YTVVLLLFMRKRSRFQSNVQR
		<u> </u>				KLMTNF
19172	49540	Α	19281	10	426	
19173	49541	Α	19282	1	900	
19174	49542	Α	19283	314	451	AGLFKSRKVSSICIRFE*ILTSCL GVRISTGKLVARTHGSFADAE
19175	49543	Α.	19284	1	1139	MWGIGSRTPIYNKILVYSSPEVS
					}	CEKPKYVKSQTSTHTGFESHKR
					,	YIFYLRLVGKNLHVSAMGLGK
						SQERRKKALKSQPQQPADSFPS
		:				LPTVEGLTTSGEVAAGFWLSGE
						ALSSC*LLSASVHLLLACLGMG
		<u> </u>				KCSCSGFWLYF
19176	49544	A	19285	1	864	
19177	49545	Α	19286	306	444	AGLFKSRKVSSICIRFE*ILTSCL
	<u> </u>	ļ				GVRISTGKLVARTHGSFADAE
19178	49546	Α	19287	189	1089	HHTAHCQ*CWL*S/DHGINPLRS
						WVGTG*V/GVRGKVQYADLGA
1						ENWKPISNLHDMSSSHSKTLGY
		1				KRLTKSNPISCQILLYKSRSKGR
						KNQRSTRTHCHHPSPKIYSASA
		l				KEPWVLATNLPVEIRTPKQLVN IYSKRMQIEETFRDLKSPAYGL
		i				GLRHSRTSSSERFDIMLLIALML
						QLTCWLAGVHAQKQGWDKHF
						QANTVRNRNGSLLTNWPSVPP
						TIKEEENSEEELAAATTSKEQEP
						IGTOLDAVRTPEPLEEFPKREDQ
						EGSPPETSLPYKWVVEAANLLI
						PAVGSSLSEALDLIES
19179	49547	A	19288	1	549	
19180	49548	Α	19289	80	430	RWLATPYLTVKLSLLPNLAVTC
						QPKREQNITSNESTDC*VIVTST
						KSDSLYTVGMLALSVRAIRCPL
		1				YLLTGLISVSKNGLWYCELQSH
						YTVVLLLFMRKRSRFQSNVQR
						KLMTNF
19181	49549	Α	19290	80	430	RWLATHYLTVKLSLLPNLAVT
						CQPKREQNITSNESTDC*VIVTS
						TKSDSLYTVGMLALSVRAIRCP
						LYLLTGLIFVSKNDLWYCELQS
						HYTVVLLLFMRKRSRFQSNVQ
			L_			RKLMTNF
19182	49550	A	19291	773	1015	AGLFKSRKVSSICIRFE*ILTSCL
		1				GVRISTGKLVARTHGSFADASF
						STDLYHGTLKPASLTMSGVVLL
		1				GRMLARSARNWS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19183	49551	В	19292	1	1488	
19184	49552	A	19293	2	133	YAQTDDNRRNLILNGTPLRRVK VLVFDIEYP**KVYGYNSTLA
19185	49553	В	19294	1	879	
19186	49554	Α	19295	80	428	RWLATHYLTVKLSLLPNLAVT CQPKREQNITSNESTDC*VIVTS TKSDSLYTVGMLALSVRAIRCP LYLLTGLIFVSKNDLWYCELQS HYTVVLLLFMRKRSRFQSNVQ
			1000		1,700	RKLMTNF
19187	49555	В	19296	1	1575	
19188	49556	С	19297	1	2187	
19189	49557	A	19298	473	613	AGLFKSRKVSSICIRFE*ILTSCL GVRISTGKLVARIHGSFADADT
19190	49558	Α	19299	410	547	AGLFKSRKVSSICIRFE*ILTSCL GVRISTGKLVARIHGSFADAE
19191	49559	Α	19300	876	1005	
19192	49560	A	19301	167	263	KLIEAIWSS*FITGSSLRHNERLG ASTSSGEY
19193	49561	Α	19302	258	396	AGLFKSRKVSSICIRFE*ILTSCL GVRISTGKLVARTHGSFADAE
19194	49562	С	19303	1	1641	
19195	49563	Α	19304	1221	1327	ALHPRNRHLHKERLGCIPVAC* LYLFGQYDAHCTC
19196	49564	С	19305	1	1353	
19197	49565	В	19306	1	2047	
19198	49566	A	19307	80	429	RWLATHYLTVKLSLLPNLAVT CQPKREQNITSNESTDC*VIVTS TKSDSLYTVGMLALSVRAIRCP LYLLTGLIFVSKNDLWYCELQS HYTVVLLLFMRKRSRFQSNVQ RKLMTNF

1	1		l .		Amino acid sequence ( X=Unknown,
of peptide	hod	in USSN	i .		*=Stop codon, /=possible nucleotide
sequence		09/540,217		or peptide sequence	deletion, \=possible nucleotide insertion)
49567	Α	19308	1	2597	MAMFMLLAAALLSTFLTNDVA
					LYTVVPLTITLKRLCEIPYNRLII
		i			FEGAGSQRWFATDANWQPAK
					YSYLGTFWSFVCRIYCPNGTAG
					WRNDADAPAPVLAMQYHTGV
		•			QTPEWKPRLVWSCLGLYIVFLT
					ALEFKQELWGLVIVAAGFALL
					ARRVVLSVDWTLLLVFMAMFI
				٠	DVHLLTQLPALQGVLGNVISNV
					PSTILLLNYVPPSLLLVWAVNV
					GGFGFLTCRHARIAAPADLKDG
					LAVHEKRETRSLTEKTGKMLLP
					KLGVGSRECGGNVKEVGIRAIE
					EFGPYKINGDKEIMRRMDDLLQ
					GFDAQHRMKLPAQPTSLATKS
					ARRVRYPLFCPSIFNGFFEEFIFH
					RFLAEQALEFFNLLHGGSKFRG
					RNNLFSGGDSSRKNQRSTRTHC
					HHPSPKIYSASAKEPWVLATNL
					PVEIRTPKQLVNIYSKRMQIEET
	ĺ				FRDLKSPAYGLGLRHSRTSSSE
					RFDIMLLIALMLQLTCWLAGV
					HAQKQGWDKHFQANTVRNRN
					CKMATLKEKLIAPVAEEEATVP
					NNKITVVGVGQVGMACAISILG
					KSRAMEHTTAGAPYTPRQQTC
					QCNMMSLADELALVDVLEDKL
					KGEMMDLQHGSLFLQTPKIVA
					DKDYSVTANSKIVVVTAGVRQ
					QEGESRLNLVQRNVNVFKFIIP
					QIVKYSPDCII\IVVSNPRDI\LTY
					GTWKLSGLPKHRVIGSGCNLDS
49568	A	19309	435	777	RWLATHYLTVKLSLLPNLAVT
	-				CQPKREQNITSNESTDC*VIVTS
					TKSDSLYTVGMLALSVRAIRCP
					CRLVDWADIPEQKQLMVLRAS
					VALHGRSVTLYEKAFPAFRAM
					FKRSS
	of peptide sequence	of peptide sequence hod sequence 49567 A	of peptide sequence hod in USSN 09/540,217	sequence   09/540,217   codon for peptide sequence   49567   A   19308   1	of peptide sequence hod grade se

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19201	49569	ĪA	19310	69	1270	LCLLRHDCYRGLQSLLLILCKM
		1				ATLKEKLIAPVAE\DEA\TVPNN
		1				KIT\VVGV\GQVGMA\CAISIPGK
						VSWPDEL\ALVGCFWEDKLKG
						GNDGSFQHGELIFFQTP*NCGQ
						DKGLFL*PANS*RFVSW*LAGS
						PVQPRRGRIRLQSWVPEKWLM
						VFQIPLFLKIVQVQSLICIIIGGFN
						PQLDILTYVT\WETKVGLPKHR
		ŀ				VIGSGCN\LDS\ARFRYLMAEK
						TWHFIPSSC\HGWNFGGNMGDS
						SV\AVWNGV\NVAGVSL\QEL\N
l						
l		ŀ				PENGNLTNDKLKIWKEGA*IW
l						WLKSA\YEVI\KLKGYPQLGAI
l						WI*VLAESYWNPMLKKSYPRI\
İ						HPGCPTNG*RQGMYGHWRME
		1				SPSLEPSLCIPQCPRGLNPAVIQP
						RSLKDDEVASASRKSARIPLWD
		1				NPEWTLKRPLT
19202	49570	Α	19311	52	406	HVEAAGSR*CAAPAGALLRFLG
						TPSFEDVSIPEKPKLRFIERAPLV
						PKVRREPKTLSDIRGPSTEATEF
į				İ		TEGNFAILALGGGYLHWGHFE
						MMRLTINRSM\DPKNMFAIWR
						VPAPFQAH
19203	49571	A	19312	1	783	
19204	49572	Α	19313	15	397	LLAEWTMGHTPVGGFRGAISL
				Ī		DAAP*EQGTMLVSWLCDLFLV
				İ		LRILVGKTVSSKNIFIERVNGDG
				ľ		KHENIPMSSPPRSWFPSEKRFTN
						RRKTSSHLQNLVLGAVSLRQH
						WRGGVPRNLTRAPAGAAH
19205	49573	В	19314	127	181	
19206	49574	Α	19315	242	1067	CAAPCGCPCQIPGALLPASAGV
ĺ						KTLLPVPSFEDVSIPEKPKA*DL
ĺ						LERAPTCAQK*RREP*KFKVTIR
ĺ						GP\SH*KPTEFYRKAIFAILGIGV
ĺ						VGYLHWG\HF*K*WRLTIQPAL
ĺ			1			WTPRTWFAIWR\VPAPFKPITRK
ĺ		1		İ		SVGASHGGEAKGAI*HYLTP\V
İ	}					KAGRLV\VEMGGRCEFARSAK
ĺ						VSLTQVALRSCPFASKRL*ACG
						TPREGCEQDQEERERQQPRTPW
ĺ						TI*AE*PTAQHGWAYGKVPEPH
	-					MTLTPQGEILGARFYMPKTCVV
i	I	ı	1	1	1	1
Į		F	į.	i		SVGDILYIGY

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	1	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19208	49576	A	19317	104	763	SAAADGPDRLRRRPPANLRARP
19200	7,570	^	17517	1104	1703	VLAPASATASARRACSPPHPQT
						MSESK\SCPEYASFFAVMGASA
						A\MVFSALGAAYGTAK\SGTGI
						AAMSVMRPEQIMKSIIPVVM\A
						GIIAMLRPGWVAVLIANSLNDD
						ISLYKS\FLQLGAGLSVGLSGLA
						QPAFA\IGFVG\DAGVR\GTAMH
						AADIFRGA*FLILIF\AEVLRL\Y
						GLIV\ALILFHKVDLFRAHQPTE
19209	49577	A	19321	36	164	GET V VIEWET THE V EEK TO THE TE
19210	49578	A	19322	190	743	NDRGKEGSLSTVPSIRNTDSGH
17210	17270	`	17322	150	, 13	LAPRRKPQLLKPSLNTSFPKAR
			ĺ			KIHKGEHEVASAGGNPPAPILP
İ						KVPP*FNIHRRGPPWPDHVRFG
				1		SLF\NTLFL\NW\CCLGVSLAFRL
						LP*KSRDTGKMGLGDVEPGAQ
						ALCLPPPKCLNI\WALILG\IL\MT
						IGFIL\LLVFGSVDSLTILWFQIIQ
						GKTGGY
19211	49579	A	19323	2	257	SK1001
19212	49580	A	19324	<del> </del>	1227	MGRARKSGLGQRSRPTASRSEA
19212	17,500		17324	1	1227	AVQPGVRKARGAGNWRVGLQ
				}		TGEAAPSPHRDLRDTPDPRPWL
						ARTHRMTTTLVSATIFDLSEVL
ľ						CKGNKMLNYSAPSAGGCLLDR
:						KAVGTPAGGGFPRRHSVTLPSS
			İ			KFHQNQLLSSLKGEPAPALSSR
						DSRFRDRSFSEGGERLLPTQKQ
						PGGGQVNSSRYKT\ELCRPFEE
		1				NGACKYGDKCQFAHGIHEL\RR
1			1			LTLHTKD*TELCRTFHTIGFCPY
						GPRCHFIHNAEERRALAGARDL
						SADRPRLQHSFSFAGFPSAAAT
						AAATGLLDSPTSITPPP\ILSADD
		ł				LLGSPTLPDGTNNPFAFSSQELA
		1				SLFAPSMGLPGGGSPTTFLFRP
		1				MSESPHMFDSPPSPQDSLSD\QE
		1				GYLSSSSSSHSGSDSPTLDNSRR
						LPIFSRLSISDD
19213	49581	A	19325	204	456	
19214	49582	A	19326	1	323	IVEYHGPGVDSISCTGMATICN
17214	17.502	` `	17520	ľ		MGAEIGATTSVFPYNHRMKKY
		1				LSKTGREDIANLADEFK\DHLVP
						DPGCHYDQLIEINLSEMV*PDM
						LPAPRLAHGVTVVVVGGFLK
19215	49583	Α	19327	2	131	E. H. C. L. L. L. L. L. L. L. L. L. L. L. L. L.
19216	49584	C	19328	2	283	
19217	49585	A	19329	1	206	
19217	49586	В	19330	217	435	
17210	77300	12	17550	1217	1.55	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19219	49587	Α	19331	1	565	PWE*SPKGIGVSCGPLSGWS/SP
						KD/VILKVQ/GILRFKGGTVQIV
						ESTGLG*TPFSCTGMRQFCNMG
						AEIGATTSV\FPYNHRMKKYLS
		ŀ				KTGREDIANLADEFKDHLVPDP
			:			GCHYDQLIEINLSELKPHINGPF
			:			TPDLAHPVAEVGKVAEKEGWP
						LDIRVDALACQSLAFGAAWVQ
	l					CGCEEHGAWSQMAWA
19220	49588	Α	19332	117	2603	GQHCLLPWGVFPFLKKALGV\R
		i				Q*\HVASVLCQRAR\VAMSHFE
						PNEYIHYDLLEKNINIVRKRLNR
		·				PL\TLSE\KIVYGHLDDPASQEIE
						RGKSYLRLRP\DRVAMQDATA
						Q\MAMLQFISSGLSKVAVP\STI
					}	HCDHLK\EPQVGGEKDL\RRAK
		l				DINQEVYNFLATAGAKYGVGF
						WKPGSGIIHQIILENYCVPWLSF
			E			*LGPDFHTPNGGG\LGGI\CIGSC
						GWPNAWEWNGLGIPWE\LKCP\
						QVIGVKLTGSLSGWSSPKDVIL
						KVAGILTVKGGTGAIVEYHGPG
						VDSISCTGMATICNMGAEIGAT
						TSVFPYNHRMKKYLSKTGRE\D
						IANLA\DEFKVHLVP\DPG\CHY
						EPTELKINLSEL\KPHIQWGPFTP
						DRGSPCWQEVGKVGREGKGW
						PL\DIRGGS*LVACTNFKLWKIW
						GRSAA\VAKQALAHG\LKCKSQ
						F\TITPGFPRQI\RATFE\RDGYAQ
						ILRDLGG\IVL\ANACGPCIGQW
						D\RKDI\KMGEKNTI\VTSYNRN
						FTGRN\DPNPETHAFVTSPEIVT
		1				ALAIAGTLKF\NPETD*PDGAR
						MGKKFQAWRFPD\AD*VFPKG
						EF*PRGRDTYQHPPK\DSSG\QH
						VGT*APRSQRLQLLDPFDKWD
						GKDLEDLQILIKVKGKCTTDHIS
						AAGPWLK\FRGHLDNIPN/NHLL
						IGAINIENGKANSVG\NAVTQEF
						GPAP*QLAR\YYQETLASRLGW
19221	49589	В	19333	1	1311	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	1	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19222	49590	Α	19334	2	1390	PRVRPRVRARPTKMSALGSLSL
	ŀ	1				PSVLWGQLF\AVDSGNDVTDIA
ŀ						DD\GCPKPPEIAHGYV\EHSVRY
						QCKNYYKLRTEGDGVYTL/NTI
						EKQWIKLRLFGD\KLP*IVEADD
:						GCPKPPEIAHGYVEHSVRYQCK
						NYYKLRTEGDGVYTLNNEKQ
						WINKAVGDKLPECEAVCG\KPK
						NP\ANPV\QRDPGVGHL\DCQRS
						FFPWARLKDGFPHH*SSTHQGP
		l				RWIQLNQWLA*PRLKNLLSLN
						H\SENCQQRKDICPYF*HFMWG
						KKQLVGD*GRLFLQP*LTPRVD
						IGLI\KLQTEGCSC*LRRVMPICL
						PSKGLWQK*GRVGLCFLAGGR
						NAQF*NLLTHLKVCPCCPVA*P
						KTQCHKAIYERPSNKSPRKRKN
						TRKKPL*GVQAPILK*NTPFLCL
						AMS*RYPKERHPAYG/QNAGAS
						ALCPFHRPLEEDTPGMPTGIL\S
		ŀ				FDKS\CAVA\EYGVYV\KVTSIQ\
						DWVQKTIA
19223	49591	Α	19335	65	665	GVRGFWAGTMASRAGPRAAG
						TDGSDFQHRERVAMHYQMSVT
						LKYEIKKLIYVHLVIWAAAWLL
						R*AVGHPEALV/DMIRWPCPIQ
						WE/YPVFC*AFLPSLLGPSFPFPR
						NNIRLTWVL/YHESSMGIFFSIG
		İ				SIHFIGQHGDVPWLHSSFYRHG
		l				KAYRFLFGFSAVSIMYLVLVLA
						VQVHAW\SLYYSKKLLDSWVH
						QHTGEEA
19224	49592	A	19336	65	665	GVRGFWAGTMASRAGPRAAG
						TDGSDFQHRERVAMHYQMSVT
	ŀ					LKYEIKKLIYVHLVIWAAAWLL
						R*AVGHPEALV/DMIRWPCPIQ
						WE/YPVFC*AFLPSLLGPSFPFPR
						NNIRLTWVL/YHESSMGIFFSIG
						SIHFIGQHGDVPWLHSSFYRHG
						KAYRFLFGFSAVSIMYLVLVLA
						VQVHAW\SLYYSKKLLDSWVH
		L				QHTGEEA

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19225	49593	Α	19337	71	1265	RCSKSLRMSTFFSDTAWICLAV
1						PTVLCRTVFCTYKKSSGQLWS
						WMVCLAGLCAVCLLILSPFWG
1						LILFSÉSCFLMHTYLSGQELLPV
						DQKAVLPTGGDCGLDHALSKY
						LDELGFTVFAGVLNENGPGAEE
						LRRTCSPRLSVLQMDITKPVQIK
						D\AYSKVAAMLHDRGLWAAIN
						NAGVLGFP\TDGELLLMTDYQR
						CMVVNFFGTVEVTKTFLPLLR*
}						SKGRLVNVTSMGGGAPMERLA
						SYGSSKAAVTMFSSVMRLELSK
						WGIKVASIQP/GEGFLTNIAGTS
						DKW\EKL\EKDILGPPPPLRVQE
						DYGQDYILAQRNFLLLINSLAS
						KDFSPVLR\DI\QQAILAKSPFAY
						YTPGKGAYLWICLAHYLPIGIY
						DYFAKRHFGQDKPMPRALRMP NYKKKAT
19226	49594	A	19338	1	861	NIKKAI
19227	49595	A	19339	248	930	DVHQLGNMAVIHSHLWEGLQE
						KFLKGEPKVLGVVQILTALMSL
						SMGITMMCMASNTYGK*PLFP
		1				CISGYTIWGSVMFIISGSLSIAAG
					•	I*N\TKGLVRGSLGMNITSSVLA
						ASGILINTFSLAFY\YSITPYCNY
						YGQPPN*LSMGTMSHLKGVW
				,		MGMVLLLSVLEFCIAVSLSAFG
						M*SALLVPPGGGCVNSAITFFT
						WAETASSTTLNEGLRGTQRFNR
						QMLPEIYA
19228	49596	A	19340	167	1164	RNLRLPSSALAGVAPSHREPRV
						RTISQTDRRKDA/PDPARRDRVE
						LAPAYTSPPRCSANVGTSTRRL
						RALCPARVRAPLRLPPPASSPLG
						SEEQRRKVPAAARMRPRAGAG
						RSTLPWLPAT/WPGLPSLPLYL*
						ICGGNNSY*R*RKGEQFE*RNSS
						*DYRYLPSSPASPNV\RERQPNE
						NGPDDKEKDRPERPSPLDTSPL
						LDPPAVSDIAGKMSDEFSLADA LPEHSPAKTSASYPDHMVLPTD
						PAAAGPLGPWGSMSSGPWVPG
						MGGQYPTPNMPYPSPGPYPVPP
						PPQASGAAPPVPWGTVPLGAW
						GPPAPYPAPTGSYPTPGLYPTPS
						NPF
L	L	LJ				171.4

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19229	49597	Α	19341	2	716	SDPSPLLDPTAVSDIAGKMSDE
						FSLADALPEHSPAKTSAVSNTK
						PGQPPQGWPGSNPWNNPSAPSS
						VPSGLPPSATPSTVPFGPAPTGM
						YPSVPPTGPPPGPPAPFPPSGPSC
		1				PPPGGPYPAPTVPGPGPTGPYPT
						PNMPFPELPRPYGAPTDPAAAG
						PLGPWGSMSSGPWAPGMGGQ
						YPTPNMPYPSPGPYPAPPPPQAP
						GAST/ILFHGAPFPPGAWGPPAP
						YPAPTGSYPTPGTLS*FPPGAW
10000	10500	<u></u>	10242	5.4	170	GPPAPYPAPTGSYPTPGTLS
19230 19231	49598 49599	B A	19342 19343	3	178 249	RHKGSPSPHQTQEPSWLHPVDP
19231	49399	A	19343	ادا	249	ALGPQVELPASLVLCARTPQPL
	1	l				GGR*DWV/PGTPAGPQESRAAP
						VPASAPPSTPRGRGGRI
19232	49600	A	19344	2	504	WAPVRPEPPRRAPPPAPQRPVP
17232	147000	l'`	17311	-		STTQGLRSAGA/PAPGLAGSYP
						CSPGYVNAPIDSLYLATFMGIR
:						VKRPPNR/PLCEQ*SFYHLGAGE
						*PGKSRVPIPQFQSP\GDPTVRV
						V\PRR*SVPPTSCATPRSWCCSD
		1				T\PGHPLGSPEQLPQPGPAAELP
		1				GGVVPRSTPPCPAL
19233	49601	A	19345	3	335	
19234	49602	Α	19346	2	447	CACTPQPLGGRWDWAPWSRG
						RRSSGRLRLHRNPRRRGKAQA
						WRAAVPRPAPREGS*GPGRPAA
						PSAGPPSPRPPGTPAGPQAPRAA
						PVPA\PRLPPHLPAS*GSGLRPSP
				ĺ		AQKGAPTVQ/PVG*RAPQVPPK
						WEPRQRRRRERARAVRTA
19235	49603	A	19347	108	729	QHAGSPQSPRLLSAPPLPGLPL
						WWHLRSPSAPPLHCGSPFLGWP
						RLEPTPSACREVWRERHKWEP
]						G/PACGACGPAGVPGGRG/PWW
						APHSEHPASPAGPG/PMGNLAP
		-				GPVAAEGVLSPPAVPAHRRCAR
						FLAGP*L/PFPRGRAWDLQPAM
						PEPPT\PPWAPV/PAEPP*RAPPP
		1				APQRPVPS\PPKGAQDWQAAPP
		1	<u> </u>	l		AAPVRDPLGEASWAPE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19236	49604	A	19348	11	816	MWESLTLSRDFLNGFDQNADS
					}	GIDNKVQVEVASDEDEELVGN
						WNKERENPFSGEKLKPAAEICIS
						YKEPNANHQDNGENVARACQ
		1				RPLQHPLPSQAQRPRRKQRFPG
						PGSGPTWDLVPCIPGAP/RLD*K
						GPRKPKSWYSEGQRYESPSFGIP
						LGSDLF\PTNALLLKPYKLPPSR
						PRDYCGRAAFPRGRVRDLQPA
						MPEPPT\PPWAPVRPEPPR*APPP
						AP/PAPSPIDHPRAEECEHTAQD
						WQAAPPAAPVRDPLGEASWAP
		<u> </u>				ESGGDVESLYV
19237	49605	В	19349	4	669	
19238	49606	В	19350	1	930	
19239	49607	A	19351	170	444	
19240	49608	В	19352	11600	1122	NOT BUILD COUNCIL BEGINNERS OF
19241	49609	Α	19353	1628	3591	KGLRLKLCSFVQLRFSIRNFRLG
ı			<b>!</b> }			ILTYWKKEFEIVGWPRLEPTPL
ı			:			ACREVWRERREREPGLRAALA
ı		1				GQLEFRVGVGLAGPSLGAAGQ
ı						PCWPRVIRDLAPGPVAAEGVLG
ı		1			Ì	PPAVPAHRRCAGFLTEP*LPSRG
ı						AGLGTCSPPCLSLPPTPWAPVRP
ı		1			}	KPPQRAPPPAP/PAPSPIDHPRAE
ı						ECERRAHDWQTAPPAALVPDP LGEASWAPESGISDEYITPMFSF
ı	ŀ					YKSIGELKMTQEEYALLTAIVIL
ı						SPAWATERHPVFKKRERKKPSN
ı						CLEGVIKAAAELAVGLPWNLG
ı						KFHTGDGTSAGLQGIAPNSVGP
ı						GQPAPEWMWGWERLLTDQDK
ı		}				TQGSAFSCTATFQDAQKQLLQ
ı		1				ARFIHLSGLWEQQDGLWLSNW
ı						LSLKGQIGGPALTEPLRLSVNR
ı						GTWPPKLRDARRRSAGRADRR
ı						APTGRVLPGSPGAGPQTTPPPY
ı						DWSLRNRRRPAATDPEAQAAA
ı						AAAAVGQGRVGPAQIEPSCRPA
ı						GRSDPPAPGPEAEVCWVSRLRE
ı						PVRSAFAWLRQSRALIGSWGR
İ						QGDVAPAFTCPPNAERGATGV
ı		1				RGGWGCWCGRAQLGRRDPAA
İ						TRPALNPERYQSPKSMQGLERP
ı					1	HVRNWGIELETLRDAFRPQPSV
İ						WRTARGEAGKRKGRLVECLGS
ı		1				l i
ļ.			Į.	B .	1	QRKVLEGVLLYAVTRGSFPNLF

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
19243	49611	A	19355	951	1453	FQTQEPSWLHLVDPTPGLQVEL PASPVPCAHIPQPLGGRWDWVP
						WSRGWCSSGRLGPHRSPWSGW
						EAQAWQAAGPEPCPMGRQVRP
	-					GEKSSAVPVGQPG/SGHSVHPP
ŀ	1		1			QPLARVLSPPLPGASRAGRLLR
		}				VWGPPSPRPPGTPAGPQARHTA
						PVPARASPSTPPCKLR
19244	49612	Α	19356	111	658	LPSRGAGLGTCSPPCLSLPPTPW
						APVRPEPPQGAPPPAP/PAPSPID
						HPRAEECERTARDWQAAPPAA
	1					LVRDPLGEASWAPESGGEPTPS
						GPGVAGSGSTHTHALTHACTH
						AHTRAEPLRLPPPPRPLPPPLSPP
		1				/PPAPLLPSKPFSLSTDRISSKSR
						QQKVPFGPRDPEPRGVRAGSW
						NRTTH
19245	49613	В	19357	1	552	
19246	49614	Α	19358	1299	1813	AHFPRAGTGAACSACWPAGVP
						GGRGLGGPRTRSSRPCWPRAK
						GDLAPGPAAAEGVLGPAAVPA
		1				HRRCARFLAGP*LPYHGAGLGT
ŀ						CSPPCLSLPPTP/SGSCAPEPPRR
						APPPAPRHPVPSTTQGPPWETC
						LLSIYGNAMNDERAWSLEGFW
10047	10.61.5	ļ	10250	380	905	VLKPSDSWMTSLSERTISFF
19247	49615	Α	19359	380	1905	WIPHWGCRWSCLPVLRRALAF
						LSPWVVDGTGRRGVGSGAGRG
						GSGRTGAHGVGGRLRHGGLQV PSPAPREGS*GPARNRAQ\PGGP
			,			ALLGDSVHPPQPLAQVLSPP\FP
						GPAGLAGCSECRGPPSPRPPGTP
İ						AGPQAPHAAPVPARASPSTPPL
						VHLTNDVLRVKVKWAFFFTLH
19248	49616	A	19360	1	1052	MGLAERCPLNHLTLTVSSVKCS
17210	1,5010		17300	1	1032	FLLAHILSVAGACYGVLRGAPH
1		1				TFPGITCPGCSQENNPIKKWAE
		l				DLNRHLIQEDTQMANKHLKDA
	ĺ					PHRYIIRELOVKPTMRYCHAPT
						GVAHIQSTNTPNAGEDVEHQEL
				}		TFIADGSPVTTRGHMDLVQTSL
						SRTLGVRSTGSWSACAEFSHGY
						CLAKRLLQLLKHGNGNSNSGG
						NAECITRHCQKPLTGILSFILAL
						NKVKNTRPFDRCHSEGSEKEPS
						FCVTCPRRPEVNQPSPYMPQPG
						KSSTHLSVVQLRALEVGEPQAF
						PPVE/PEGPAGLRHPLWL*V*GQ
	1			ļ		GSGP/SSPPCLSLPPTPWAPVWP
		1				EPPRRAPPPAPQRPVP\RPPKG
L	<del></del>	ــــــــــــــــــــــــــــــــــــــ	l		<u> </u>	

SEQ ID NO:	of peptide	Met hod	SEQ ID NO: in USSN	location of first	codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19249	49617	Α	19361	417	1092	KWCSKRTAFSFSLQGGVEGEA
						RAGTRAACGACGPAGVAGRRG
}						LGGPPHSGQPASPAGPG/PVRDL
						APGPVAAEGVLGPPAVP/APPA
						LCSIS/PPGCSCLPAG/PRLRTCSP
						PCLSLPPPPWAPVRPKPPR*ATP
						PAPRRPVPST/NPRAEECGRTAR
						DWQAAPPAAPVRDPLGEASWA
	1					PESGGDVENLYVQLRDCKYTN
						QHPVSSSGFVNAPISTLYLVNL
						VGTWRIFMSS
19250	49618	В	19362	817	1596	
19251	49619	A	19363	350	644	CDACEDDADE A CODED CD DACED
19252	49620	A	19364	60	1362	SRVEPRVRSASQDSRSRDNGPD
						G\MEPEGVIESNWNEIADS\FDD
						MNLSESL\LRGIYAYGFEKPSAI QQRAILPCIKGYDVIA\QPQSGT\
						GKTATFAISILQ\QIELDLK\ATQ
1						GLGS*APT/RENLAQQIQ\KVVM
İ					}	ALG\DYMG\AS\CHACIRGAPTC
						VAEVQKL\QMEAPHIIVGYPLG
						RVF\DMLYR\RYL\SPKYIKMFV
						LDEA\DEM\LSRGFKGQIYGHIQ
]		ŀ				KAQAAPPRVVLLSATMPFD\VL
		1				EVTKKFMRGPH/IRILVRKEELT
						L\EGIRQ\FYI\NVEREEW\RLDTL
						CDLYETPDPSPQASHLPSTTRR\
		1				KVDWPHPRRMHALRFPWESAQ
						HEDMAQKERR/DVIMKEFRLAS
						SRV\LITT\DLLA\RGIDV\QQVSL
						SHQLNY/SFPTTRENY\IHR\IGR
						GGRFG\RKGVPINMLTE/EKTKR
						NLEDIETF\YNTSI\EEMPLNVA
19253	49621	Α	19365	911	1278	NCGIVISKHLVFTLA*MMCHFL
						ASLQRKGILKRSASV*AHSH/DP
						SFHMKIYNAGW\GHRT*HPAGG
						FAGPPPWGSVWGLAARTRENA
						PRPALPLPRPAPSGLAHGADGD
	į					SNTEKIQSRDNRKG
19254	49622	Α	19366	26	743	ETSADMAPSVPAAE\PEYPKGIR
						AVLLGPPGAGKGTQAPRLAEN
						FCVCHLATGDMLRAMVASGSE
						LGKKLKATMDAGKLVSDEMV
						VELIEKNLETPLCKNGFLLDGFP
						RTVRHAEMLDDLM\EKRKEKL\
						DSVDLNFSIPDSLLIRRITGRLIH
						PKSGRSYHEEFNPPKEPMKDDI
						TGEPLIRRSDDNEKALKIRLQA
						YHTQTTPLIEYYRKRGIHSAIDA
						SQTPDVVFASILAAFSKATS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				<u> </u>		
19255	49623	A	19367	40	1002	VASGTRTELQDRRFRDTSGDM
		1				FPSVPP*EPEYTKGIRTVLLWPP
						GAGKGTQ/SGETEV*MACSLGY
						IASNLQSLHLNDSKAPALSAAIY
						CPRHPNWAENFCVCHLATGDM
						WR\AMVASG/SSELGKKLEGNL
						WDGWGNWVE*WKWVSWSSF
		l				GEGILENPLVAKNGFLL\DGFPS
						GPVKARQK\SPMDLMGEGGKE
			1			KLDFC*FEFSIP\DSLL\IRRNHKE
						GLIPPPKKWAGSYPRREFQPLP
		l				KEPP*KMNITGGNPLIRRSR**M
						KRPLKIRPASPTHTQT\TPLIE\Y
		Ì		1		YRKRGIHSTI\DASQTPDVVFASI
1				1		LAAFSKATCKDLVMFI
19256	49624	Α	19368	793	1152	SGRKRRGCGGQSVKQAPALTP
į						GPPAPVFPIPEAPAACAPHHPGQ
						GNRRKCNFPPVCPT\PKTASSCP/
						EPLPPPPSLQR*PCPCHPWHHLE
						FSFLDEVTFSFNKTFPKRLVSSK
						MLPLKVGH
19257	49625	A	19369	1	873	
19258	49626	В	19371	1	322	
19259	49627	В	19372	64	2250	
19260	49628	Α	19373	20	368	CEARGGSGHGGRAERAGVCGG
						PAELLPKGSKEEQRDYVFYLAV
						GNYRLKEYEKALKYVRGLLQT
	1	1				EPQNNQAKELERLIDKAMKKD
		į				GLVGM/AIVGGM/ALGVAGLA
L						GLIGLAVSKSKS
19261	49629	Α	19374	3	492	
19262	49630	Α	19375	62	570	GSRGQWPSEAVLNELVSVEDL
						LKFEKKFQSEKAAGSVSKSTQF
	į					EYAWCL\VRSKYNDDI\RKGIVL
					,	L\EELL\PKGSK\EE\QRGFTSFYL
						ARGGGTTRLQGITEKALKLRSA
				,		GLLAAQEPQEQPRPKETGNGLI
						DKA/L*KKDGLRGAWPIVGGM
		1				ALGVLAGTCRSSSDFAVFQ

SEQ ID		1	SEQ ID NO:	1		Amino acid sequence ( X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	Sequence		027510,217	sequence	or pepme ouque	positive nucleonal insertion)
19263	49631	Α	19376	1	944	MACHLQAGSPGNLVVYFQSKP
			•			EGLRTREPMMSVPVQVKGPST
						RKLMMSEDPLTLGNWHELRVS
						RTAKNGILQVDKQKIVEGMAE
						LQQVVGKNMDSGVGPVVHGG
						GLTRTGYLLNPESLQQKAFERR
ĺ		}				QIQGCDEKLACVGKQTLGTWP
	İ	İ				LSESPSLTRLYCRRQLTQGSLR
						MSLNNAGVPGQTAPDQRSGEG
						VAAPAGPGEEVGREEGEGAAA
						GKRQCPCQQGSLCVDTSRPRLP
						KSQAQIPSSSRSSPSSSEQQRGQ
						LPSVREHVEGGQGEERAPHCD
	ļ					GGGRVLLASGLAWLGTENIMD
				1		LASSPKDDWGVAVPSRGCERPS
						TFTLVLVPSPIVRRRACSGQPISP
						GGGQETHGADTSHLTNP*ICLL
						SKAFCCRDSGLSK*PVLVRPPPC
£						\PQALPQSPCSYLLLAVALPFPP
	<u> </u>	_				LSSAYPPVRFHSLLCEIHEARAS
19264	49632	Α	19377	1177	1603	WVSSFSSSSFSSFPPPPPPPPPPPP
						PPSFLPPPPPPPP/PPI*DRARRCQQ
	İ					TSIPLNKLNINKLLLLLLSPRPD
						DREKFTPQPQSARQREPIKGTA
						SRRGEGPICPTLSLYFLHVFLCF
						SCLSSQIHIYILRLSTVLVLIKNS
19265	49633	A	19378	1	896	FPFSLLPPPKMPKGKKAKGKKV
						APAPAF\VRKQDAQE*WLNPL\F
						KKRP*DFR\IGQ\DIQPQKRPPPL
		l				L*KWPR\YIRLQR\QRAILYK\RL
						K\VPSAITPVSPRALDPANQLLS
			ŀ			LLKL\AHKYRPRDKAQRSKQR\
						LLA\RAEKKAA\GKGDRPNQRD
						PPVLRARS*HPSPTLVEEQEKLS
						WVVIA\HDVD\SIELGLSFLA\AL
						CRKNGGSPYCIIKGKGQDLGR\
						LVHRKDLAPLVAFTQ\VN\SED
						KRRFWLKLVGSYQGPNYNDRY
						RYDEIRRHWGGNVLGPKSVARI
						AKLEKAKAK\ELATKLG
19266	49634	A	19379	3	491	AKEALDWEEGHQ*DQRL*SRN
						SQLGICGGLRGFRGDRAASVFS
						R*AEKESVRQL*ARVPGALGVA
						ARANKHQRRWREGDSTQTATN
						TVDRRATRGRPMEWRATRVAD
						EEDGDRRRGDNRPQARTFRPRT
						SVRRAAGTNNSEGRRPTNAGT
				L		NGGRGVDIYTVN

SEQ ID NO:	SEQ ID NO: of peptide	1	SEQ ID NO: in USSN	Nucleotide location of first		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19267	49635	A	19380	1	1501	MRRSTPGVLPAAVGGARVRRA
						VGLSGTGAEAGRAGAMVEKEE
						AGGGISEEEAAQYDRQIRLWGL
					İ	EAQKRCHRMKEALSTAPFTDK
	ļ					ETHPKVSLAQVTRSSASVMVY
						GAFEVIGOROSSAAKPRRSOSE
						SLGPEFQGLWEWLPASQHIMRS
						LKKQDGMGQGYRVSLEWEVG
						QLIFLEEKVTP\EDPGAQFLIRT\
						GSVGRNRA\EAS\LERAQNFNP
						MV/DMLKVDTEDIEKKPESFFT
						QFDAVCLTCCSRDVIVKI\DQIC
						HKNSIKFFT\GDVFGYHGYTFA
						NLGEHEFVEEKTKVAKVSQGV
						EDGPDTKRAKLDSSETTMVKN
						KVF\FCPVKE\ALDGWILDACDE
						MQMAALKRTTSDYILLQVLLK
						F/RVTDKGRDPSS\DTYE\EDSEL
						LLQI\GNDVLDSLGISPDLLPEDF
						VRYCF\SEMAPVCAVVGGILAQ
						EIVKALSSAGTLP/PTTNFFFFR
						WA*KGEWGLWECLGPQVNSG
						FGQPQRCQLQHAHLYSLFPFFM
19268	49636	Α	19381	2	411	PGTSQGVFSPHRLGGAVRSPCP
						PPQDSRQASAGGAESQ\LKQRG
						FPGE*GCRSPVSCAWPQNRMG*
						GPQHREAGAWSGE*KTPIWLK
						QERCNRPGLLVTAELAAPAPRP
						DGMAPGCGGIRCTPGRPRRLLH
						SGPPSP
19269	49637	Α	19382	165	309	
19270	49638	Α	19383	1	825	LATQPEGAGRVPSIHPQGGPEA
						ALQDLRTLMGSSRLHSGKESIG
						HNSSAGSSSQGPHHPRRKTAIL
						EKGGKQPHRMLPFSPGLGSSPE
						PANAGSPDTWLCPP/AAARVL*
						NPVKPRKGRSEPRSGWASQLPG
1	ļ			-		GDSRLPLRPGTSQGVFSP/APAG
						*AGKLVLGVLSLSLKQRGFPGE
						*GCRSPVSCAWPQNRMGCRTSP
						GPSPTSPTAAGE*GSRPRRRKPG
						RVQGLQRDGPTLTQEDALTAA
						KRCPLPVTSVPQTPETSTGAAA
						PGSQPVGAPPSR
19271	49639	A	19384	1	378	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19272	49640	A	19385	381	1653	QRPRSGTFTMGRKSLYLLIVGIL
						IAYYIYTPLPDNVEEPWRMMWI
						NAHLKTIQN\LATFVELLGLHHF
						MDSFKVVGSF\DEVPPTSDENV
i						TVTETK\FNNILV\RVYVPK\RKS
						*STKEGGLFYIPCGGWRRGSYS
						LIVYDLPSSWTVE\RPGSSLVSN
						HL\RLAT*F\HFPLQLGELYDAL
						R*FLRKKVLAKYGCEP*GESGIS
						G\DSAGRGI*LSAV\TQQLL\DDP
						\DVQDPNFKIQS*IYPALSASWM
						*VLPSYQGKFKFSYFYSKSLQ\C
						RFWSEYFTTDRSLE\KAMLSRQ\
						HVP\VESSHLFKFINWSS\LLP*E
			:			GLLKGH\VYNN\PNYGSS\ELAK
						KYPG\FLDVRAAPLLADDNKLR
						GLP\LTYVITCQYDLLRDDGLM
						YVTRLRNTGVSG*LNN\HVEG\
						GFHGAF\SFLGLKISHRLINQYIE
19273	49641	Α	19386	3	667	DAWADAWVPVSSEVAYLFGG
						CHTCGGLPVAHPMMPRGSRSR
						TSRMAP\PASRAP\QMRAAPRPA
						QVG\QPPAAAPPSAVG\SSAAAP
						RQP\GLMAQMATTAAGVAVGS
						AVGHTLGHAIT\GAFSGRK*LPE
						PARP*HHFNQEASG/TPPAQHSK
						QQRHPHLQQPCP\YEIQTVFWE
						FCPGTRGDIQAL/CEGFNE\VLK
	]					Q\CRL\ANGFGPNEEVSTFGRD
10074	10640	<u> </u>	10207	0.1	1057	WKFSSHNLS
19274	49642	Α	19387	81	1257	RHQSDQTVPRLSRPHVQSQKSS
						MAAAAKPNNLSLVVHGPGDLR
						LENYPIPEPGPNE\VLLRMHSVG
			:			ICGSDVH\YWEYLS/RFGNFIVK
						KPMVLGHEASGTV\EKVGSSVK
						AP*NQGDRVAIEPGCSPEEN*W
						NSC\KMG\RYNLSPSIFFCATPPD DGNLCRFYKHNAAFCYKLPDN
						VTFEEGALIEPLSVGIHACRRGG
						VTLG\HKVLVCGSWANRGWVT LLVAKAMGAAQVVVTDLSATR
						` '
						LSKAKEIGADLVLQISKESPQEI ARKVEGLLGCKPEVTIECTGAE
1	}					ASI\Q\AGIYATRSGGTLVLVGL GSEMTTVPLL\HAAIREV\DIKG
						VFRY\CNTWASGGFSMLCVPKS
						V\NVK\PLVTHRVSSWRKVLEA
19275	49643	Α	19388	2	431	FETFKKGLGLKIMLKCDPSDQN
17413	47043	Α	17300	2	1,21	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19276	49644	Α	19389	2	326	ARGSTRLMESRVQHL*CTRKEL
						TPEVLQEWLDELEEMMLVVH
		l				MPRFRIEDGFSLKEQLQDMGLV
			į			DLFSPEKSKLPGIVAEGRDDLY
		1				VSDAFHKAFLEVNEEGSEAAAS
19277	49645	Α	19390	2	368	
19278	49646	Α	19391	1	2123	MAQFSIVGMCLGLFNQSPTDGS
		1				GVQLFVIMNQVMAANLAETHA
		1				ALCPTLSSGTSARFRGKNQFSG
						GLPQITLSPLAQPCGRLAAMYS
			,			NVIGTVTSGKSLPSTSPLVPLLL
						SGFAPLFLKGLQNWTQTTQEG
		1				KSPSAAPASSQPQACPTDHVRE
		•				SALGACLSLPGEDGKIQKVYLL
,						SLLLIGFWDCVTCHGSPVDICT
						AKPRDIPMNPMCIYRSPEKKAT
					]	EDEGSEQKIPEATNRRVWELSK
į						ANSRFATTFYQHLADSKNDND
						NIFLSPLSISTAFAMTKLGACND
						TLQQLMEVFKFDTISEKTSDQIH
						FFFAKLNCRLYRKANKSSKLVS
						ANRLFGDKSLTFNETYQDISEL :
		Į				VYGAKLQPLDFKENAEQSRAAI
	1	1				NKWVSNKTEGRITDVIPSEAINE
	1			ļ		LTVLVLVNTIYFKVLRMALERP
						QGLPLALQLTPFFFKWRDRSPE
	1			ļ		RANGLPKATQGLWKSKFSPEN
						TRKELFYKADGESCSASMMYQ
						EGKFRYRRVAEGTQVLELPFKG
		1		ŀ		DDITMVLILPKPEKSLAKVEKE
		1			į	LTPEVLQEWLDELEEMMLVVH
						MPRFRI\EDGFS\LKEQLQDMGL
		]				VDLFSPEKSKLPGIVAEGRDDL
		1				YVSDAFHKAFLEVNEEGSEAA
						ASTAVVIAGRSLNPNRVTFKAN
						RPFLVFIREVPLNTIIFMGRVAN
						PCVNLSEALAVLVVNLTTLMK
						RTHGESLDFMERKRRPAESILL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19279	49647	Α	19392	9	1630	PALCPTLSSGTSARFRGKNQFS
1,72,7	1.5017	l'`	1,,,,,			GGLPQITLSPLAQPCGRLAAMY
						SNVIGTVTSGKRKVYLLSLLLD
						SFGDCVTCHG\SPVDICTAKPRD
						IPMNPM\CIYRSPEKKATEDEGS
				ł		EQKIPEATNNRRVW\ELSKANF
						PLLLTTFLFSTWADSKD*H*LTF
						FCSPLSIFQGFLLWTKV\GACND
						TLQQLMEVFKF\DTIS\EKTF*SR
						SHFFFA\KLN\CRL\YRKANKSSK
						LVSANRLFGDKSLTFNETYQ\DI
						SELVYGAKLQAPGTSRENAEQ\
						SR\AAINKW\VSNKTEGRIHRCS
						FPSGRPFNELTVLGGGFNTIYFQ
						GACWK\SKFSPENTRKELFYKA
						DG\ESC\SASMDVTREGKFRYSG
						AWLEGT/QVLVVCPFKGDDIT\
						MVLILPKP*EGAWAKVEK\ELT
						PEVLAKSGWD*FWREMMLVV
						HMPRF\RI\EDGLQV*REQPAKN
	İ					MGPLSD\LFSP\RKSPQTPPGIVA
	:					EG\RDDPPMSSDAIPIRHFLEVN
						EEGSEAACKYPLL*LPGRS\LNP
						NRVTFQGQQAFPWVFIREVPL\
10000	12612	<u> </u>	10000		1505	NTIYLHGAELANPCV
19280	49648	A	19393	3	737	
19281	49649	A	19394	1	705	
19282	49650	A	19395	2	757	
19283	49651	A	19396	1	1764	MANUAL I I TO DECOMA OG
19284	49652	Α	19397	1	710	MAWALLLLTLLTQDTGSWAQS
						ALTQPASVSGSPGQSITISCTGT
						NNDVGSYNLVSWYQQHPGKA
						PKIMIYEVSKRPSGVSNRFSGSK
						SGNTASLTISGLQAEDEADYYC
						CSYTSSRPD\VVFGGGTKLTVL
						GQPKAAPSVTLFPPSSEELQAN
						KATLVCLISDFYPGAVTVAWK
						ADSSPVKAGVETTTPSKQSNNK
						YAASSYLSLTPEQWKSHRSYSC
						QVTHEGSTVEKTVAPAECS
19285	49653	A	19398	1	679	
19286	49654	Α	19399	1	743	
19287	49655	Α	19400	2	736	
19288	49656	Α	19401	2	718	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19289	49657	Α	19402	1	711	MAWALLFLTLLTQGTGSWAQS
						ALTQPASVSGSPGQSITISCTGSS
						SDVGGYNYVSWYQQHPGKAP
						KLIHYEGSKRPSRVYDRFSGSM
						SANTASLTISGLQADDEADYYC
						CSFAGSSALRIFGSGTRVIVIGQ
						PKANPTVTLFPPSSEELQANKA
						TLVCLISDFYPGAVTVAWKAD
		ŀ				GSPVKAG\VETTKPSKQSNNKY
						AASSYLSLTPEQWKSHKSYSCQ
						VTHEGSTVEKTVAPTECS
19290	49658	Α	19403	3	720	
19291	49659	В	19404	1	711	
19292	49660	Α	19405	105	858	RRQDSGQSPAWPAGALLLTLLT
						HCAVPGSW\AQSVLTQPPSASG
						TPGQRVTISCSGSSSNIGINDVT
						WYQQVPGTAPKLLMYSDNHRP
1						SGVPDRFSGSKSG\TSASLGPSV
[						GLQSEDEADYYCGTWGLAA*M
	ŀ	İ				VFMSSAGGTTLTVTSQPKAAPS
						VTLFPPSSEELQANKATLVCLIS
						DFYPGAVTVAWKADSSPVKAG
						\VETTTPSKQSNNKYAASSYLSL
						TPEQW\KSHRSYSCQVTHEGST
						VEKTVAPTECS
19293	49661	В	19406	63	756	
19294	49662	Α	19407	1	757	MAWTPLLLLFLSHCTGSLSQAV
						LTQPSSLSASPGTSASLTCTLRS
						DINVGSPFIYWYQQKPGSPPQF
						LLRYKSDSDNQQGSGVPSRFSG
						SKDASANAGILLISGLQAEDEA
						DYYCSSYVGTNNFGVLFGG\GT
		ł				\KLTRP*GQPKACPPRVTL/FPHP
						PSEEASKPNKA\TLVCL\ISDFLP
						GKP*QLPWKAA*QPPSLAGVET
		ŀ				PPHPPTRSNNKYAASSYL\SLTP
						EQWKSHRSYSC\QVTHEGSTVE
		l				KTVAPTECS
19295	49663	Α	19408	1	224	YISLPAGVLSLACCTLYGISWQF
						DPCCKYQVEYDAYKL*RLPLH
1			ļ			TLTCSRPTRDNRVFRSPHSVRQ
		1				QSLGRCLA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last Amino acid sequence ( X=Unknown,			
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide		
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)		
				sequence				
19296	49664	A	19409	235	955	ESTVQVSVFCHRVPARTSSQTA		
19290	49004	^	13403	233	933	LPVLLGLLGFTVSGPLNLPEPSV		
						IVGNGELVGPQTCLHCGMEIYN		
						G\ENAQ\DQF*VTELGAGP*EGP		
		ŀ				VTSYIVIE\PTRIGDETARWITVG		
						NCLHKTAVLAGTALLLHPGLR		
	•					CTLRFIPHHISPAPLGVL\SLACC		
						NL\YGIS\WQ\FEPFACK*QVEVR		
						TPNKLSRLPLHTLTLLPPRVVLV		
				ļ		RKDDLH\RKRL\HNPIGTGPPLV		
						YCVK\KIYELYAVMIFK		
19297	49665	A	19410	1	764	MRLGIGFACPWKGLADEFDLKI		
19297	49003	^	13410	]'	704	EKQRNQGRLLGFKLQHLVPFTE		
						MEKMGGDRQKHRTAPRQAER		
						RRSGRVSFQPPLQPLDVTLLPG		
İ						ATREGNCPRASRGRRPSPISHCQ		
		1				ETQGPEGRSDCTIRGVLSSIPAA		
						RPLRKGSGFPTVCEKRPEQGQQ		
						PALGP/AGPAAVPAPAASRTQW		
						GSRCPCGAGARRGYWRGLLGL		
						CNLISPLDPGSPGSGRWGSAGA		
						LSRPADSCPPWWPRAGA*RRG		
						AAAEAGNSPSRCGASRPAEEQS		
1						GAFAGPLPSSPSQ*RAPGAEV		
19298	49666	A	19411	209	536	TRSSCVTLVC*LAMVSLRVARP		
19298	17000		17411	209	330	HGVPGTMFGKRKKRVEISAPSN		
			İ			F/EHRVHTGFDQHEQKFTGLPR		
						QWQSLIEE/SARRPKPLVDPACI		
						TSIQPGAPKEVCSPRASGTRSR		
19299	49667	A	19412	124	1983	1510. 511115 5511615 11616		
19300	49668	A	19413	424	1582	PPWAAVHGDRPHRVPGTMFGK		
13300		1.	1,,,,,			RK/KRVEISAPSNFEHRVH/TGF		
						DQHEQKFTGLPRQW/QSLIEES		
			1			ARRPKPLVDP/ACITSIQPGAPK		
		i				WGETGSWPPF/RHLPEG*HGPPI		
						PGCQREPQRVSHEQ/FRAALQL		
		İ				VVDPGDPRS/YLDNFIKIGEGST		
		j				GIVCTPPSGKLVAVKKM/DLRK		
			1			QQRRELLFNERPWQCP*HP/RR*		
		İ				EAQVPLPVLQALSV/LHAQGVI		
ļ						HRDIKSDSIL/LTHD/GQVSKEVP		
						RRKSLVGTPY/WMAPELISRLP		
						YGPEVSPGWLGCPAVDSVRCH		
						FQLQRWREMGPSLGSHCQNGA		
1						LDKEEGQRPSHTGSLQWGSLG		
						VERTLRVNMYECVHACNSFYL		
						VNKLFQKDSTQEASVTMVPDP		
						DKDGKNRASQKRKAHSQGARR		
L	<u> </u>		1		<u> </u>	DI DOMINIO QUINTINO CONTINU		

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19301	49669	A	19414	488	707	TCGEGLSWTRPLPTASSLQV\H* SLSAGERPYLCDYPDCGKAFVQ SGQLKTHQRLHTGEKPFVCSEN GVNLVF
19302	49670	A	19415	16	454	GPAGRCPADCSEEKEGPSA/GD SGRPPSCT*PWKGLPVMAILPG EAPVPGPGWPEPDAGG\PLWR WGSPSPCAAALHRP\SGTTPPGR ALSGPGLERAKQRSPRVSPAPR AAPHGHPTHCLPIPAGPRPAKD HPAYGPPASCHHHLAAS
19303	49671	A	19416	166	1191	TAGSLVLAARGT*GPPPRARK* GPAGRCPADCSEEKEGPSAETP GTTPPGRALSGPGLERAKQRSP RVSPAPRAAPHGHPTHCLPIPA GPRPAKDHP\PTHQKPPPRTEGS TGGHCCGPPIFLIPNTSMLTVYP SGPVMPAGFTLSPSDTRPSLLN GQEVLRKMNGAVQYLSLCDYII PTCQCLATTVLCSVSMSLTVFA VSYMVELLLLQNAQVHQLVLQ NWMLKALPPALQDPPHVPPRV PRAARTRLPAV

2421

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19304	49672	Α	19417	1	2481	MRSGDLKFGKSAAWPSGQKEK
						LVFRGKIEEGLEYLHKKKPSAN
		i				SQDNGKEASKAFHGPLQQPLLS
						QALGPDREEWFPGQAPWPHCC
						VHPQDTAGCIPVVPAPAMAQR
						CTGTAWVTASEVAPSLDCFHIV
						LSQQVYRIHRPTSSLHPQYGKA
						TRHTTPAVSMRAAAGAEPCKA
		ļ				TGADWPKTLGAQPSHPCTMDV
						GQGFKKDDSGAALAPMAALM
		İ				GWAGIECLWLFHTEGVNCWP
		ļ				WDLGEKNGLLGQPRGPAAVCS
	į					LRTLLPASQQPQLLLRPWLKDA
						QVQLGSLLQRVQAIGLGGFYIV
:	İ	1				LSQFIDPSKASYPPWVKNSRHS
		1				TAAQPMRAAAGAEHCKATGA
		1				ELPKALGAQPSCPCVLDVGQGF
		1				KKNDFGAVGLNNWPAGFWNV
		ŀ				MGTCKSRLCFVLLSGKKSSFRV
		1				EVLTQCLDSHTLETYQFYNREV
	į.					AFEDVLESSQTGLVVQKPQTDR
						LTKRGRGEATRNLRGLVPSMP
						RSLLPSTAQPTAWKCQRGPSPY
		-				THQDMALIPSPTARCLSPAKEP
	1					KQEEVGEKSLLPDPTLPLTDPR
						LTGSTEQAHAEGLAALMSALR
						VSHLQGRGGVVTLVDSQLGVI
	1					AVSSTQFNKGPSYRLLADVQN
1						RLLPKYDSQKEAELRSWIKGFT
						GLSIRPDFQKGLKDGIILCTLVN
						KLQPGSVPKIN\ASV*NWH*LEN
•						LSNFLKAMVSYGM/NPVDLFEA

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19305	49673	A	19418	1	1299	MHQEDLRAWYLDLGLPSHQNE
1						TTDVSVVYERFSLTDATYPKQA
						KQWDFPFLRVSTAQPTAWKCQ
						RAPSPYTHQDTALIPSPTARWL
						SPEKEPKQGEVGEKSLLPDPTLP
						LTDPRLTGSTEQKKDLAAGPVG
1						SALRRRPPAVSSTQFNKGPSYR
						LLADVQNRLLPKYDSQKEAEL
						RSWIKGFTGLSIRPDFQKGLKD
		ļ				GIILCTLVNKLQPGSVPKIN\ASV
						*NWH*LENLSNFLKAMVSYGM
						/NPVDLFEANDLFESGNNMQVR
		l				VSLLALAGKAKTKGLRSGVDIR
						DKYSKKQNFDDTTMKASQCVI
						RLQITNKCASQSGMTAYGTRR
1						HLYDPKNRILPPMDNSTISLQM
1						GTNKYASQVGMTAPGTQRHIY
		l				DTNLGIDKCENSSMSLKMGYT
						QVANHSGQVFGLGRQIYEPKY
						QPGGPVAHGAPSAGNCPGPGE
19306	49674	Α	19419	135	290	HREEESKSSVFLNPRGVQGTGV
						GDHRMILGRGRAVLSKE*AAG
		<u> </u>				PRGGRHES
19307	49675	Α	19420	92	1058	KGQYLNPNRGKGGPEPPFEPPV
						PPASRSDMAQNLKDLAGRLPA
						GPRGMGTALK\LLLGAGAVAY
						GVRESVFNVE\GGHRAI\FFNRI
						GG\VQQDT\IL\AEGFHF\RIP\WF
						QYPH/IIYDIRAKTSKKSPPLQGS
		ŀ				KDLQM\VNISPAKCWSRPNAQE
		ļ				LS*ACTSRLGLGLTRER/RCLPSI
		ļ				VNEVLK/AVWVAKFNCLTA*FT
						QRAPGYSLLDSAGSLTERAKGL
						SSLIL\DDVGQSQKLSFSPRSYT
						A\AVEAK\QVAQQEAQR\AQFL\ VEKAKQ\EQRQKI\LQGRG*GR
						GLPKMLGEA\LSKNP\GYIKTSQ
		1				RFEPAPEIILQRRIAHIHRNP
19308	49676	A	19421	3	861	LDDSWEPVHPGCGDFICQWSEE
19306	49070	^	19421	]	801	SGLWSTAEVWRERRGREPGLC
		l				EGARGTARVPGGRGPPHQSGQ
		1				PVPPAPGS*GA*\PGPAAVEGVP
						GPPAVPPASAALEFSRGFSCLPR
		l				CHLVVTKIGKQEDDVSIKRKNR
						PDYWPAPEPVAGCEGSDQEQV
						KPNATLEKHFLTEGTGPRSPSC
						LSWPPSGPHAAADPAMVPETPE
						PDRPQLTKKFCEARVMAVGTS
						NVLGQTLKPSRTGGTSWSWVF
				[		YLLTANRLPFDVKRKGGGPSSI
						KPRPHYDHRLLQDFKEQVIHHF
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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19309	49677	Α	19422	808	923	
19310	49678	Α	19423	1	339	
19311	49679	Α	19424	1	621	
19312	49680	Α	19425	3	1836	MAAVEAAAEPVTVVAAVGPK
						AKDEEEEEEPLPPCEA/LRWAP
				į		VLWRGPAWATAFLEEATAEEP
						GAAPGSPPDSPDRTLRRLRAER
						RRLDS/SLLALSSDFAQVQFRLR
				:		QVVRGAPAEQQRLLRELEDFA
						LKG/IPDILGYEGPGDPASDEGD
ŀ						GLPGDRPRLRGEDQSEQEKQER
						LETQREKQKELILQLKTQLDDL
ľ						ETFAYQEGSYDSLPQSVVLERQ
		1				RVIIDELIKKLDMNLNEDISSLS
		1				TEELRQRVDAAVAQIVNPARV
						KEQLVEQLKTQIRDLEMFINFIQ
						DEVGSPLQTGGGHCECKAGGK
						TGNGCSRTGSSRTPPGNSRTKA
						EDVTKVRKTGLHLMRRALAVL
						QIFAV\TRLVVPQARSLQPCGRG
						SRLTETTLPC*R/ELEASVDRVK
						QLALTAEPHDHVITSANLQDLS
						L*GKDELTMAVRKELTVAVRD
						LLAHGLYASSPGMSLGMATIA*
	1					LLPAFSSVPEAMHPRELFVKYY
						HAKNGRAYVESPARKLSQTFA
						LPVTGGTVVTPKQSLLTAIHMV
						LTEHDPFKRSADSELKALVCM
						ALNEHRLVCWMNLICKSGSLIE
						PHYQPWSYMAHTGFESALNLL
						SRLSSLKFSLPVDLAVRQLKNIK
19313	49681	Α	19426	366	477	
19314	49682	Α	19427	3	432	WNSRDAGYEFDICFTSVHLY\AI
						RTLWTVLDAIDQMWLPVVRT
						WRLNERHYGGLTGLNKAETAA
						KHG\EAQVKIWRRSYDVPPPPM
						EPDHPFYSNISKDRRYADLTED
						QLPSCESLKDTIARALPFWNEEI
						VPQIKEGK\RVLIA

SEQ ID NO:	of peptide	Met hod	SEQ ID NO: in USSN	location of first	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	or peptide sequence	deletion, \=possible nucleotide insertion)
19315	49683	Α	19428	1	890	MRTRRTPRGEHSLSAFSSCVDK
						ISRVCKSFGLELENSIQTLWAER
				ļ.		EQSICLVAAGHLVSSSGMRNW
				ľ		CHARRHAAYKQVLIRHGEVAW
				ŀ		NLENRFGGWYDANLNPAGDEE
						AKRWEQALPDAGYEFDIC/FKR
		ł				TIRTLWTVPDAIDQMWLPV\NV
						SKDRRYADLTEDQLPSCNSLKD
						TLARALPFWNEEIVPQIKEGKW
						SDHCLGALVIPVKDGCNLHGD
						VTTVSSLTPEERALEEPLNRPTN
						A/P*HQEG*CWESHQKPVPFQT
		l				LGCLCEFSRGASRRKQARVVFR
						ERSNLCRSISGGEGF
19316	49684	Α	19429	1	443	RALPFWNEEIIP\QIKEGKRVLIA
						\AHGNSSGAIVKHLEGLSEEAIM
						VLNLPTGIPIVYELDKNLKPIKP
						MQFLGDEETVRKAMEAVAAL
					_	VCVLESLPALFHGSEMPLGPVQ
		İ	:			VYLSLISESCSSCLTLPHGSSVH
						LSITVLNPFSISV
19317	49685	Α	19430	24	1039	AVLALLRGRRGGLLLRNLLIPV
						GAASPARRQWPPYKLVLG\RHG
				1		ESAWNLENRFSGWYDADLSPA
						GHEEAKRGGQALRDAGYEFDI
						CFTSVHLY\AIRTLWTVLEMPL\
						SEMWLPVV\RTWRLNERHYWG
						S*PGLN*KQKLAAKHGLRPRLK
						I/WRRSY/D/VPPPP/MEPDHPF/Y
ł						SNISKDRRYADLTE\DQLPSC\ES
						LKDTIARALPFWNE\EIVPQIKE
						GK\RVLIAAHGNSLQGHCSRLL
						EGLF\EEAIHGSLNLPDLVFPLSI
						ELDKNFESLSSPMQFLG\DEETV RKAMEAVACPGQGPRSEGRRG
						RILSPGCTLPCPVFVPSSPSHLA
						HVTLDHICKTS
19318	49686	Α	19431	86	431	VYVSTMSLSCPASELAVRRQR
15510	77000	^	17431		731	WSSQFQDHPPGPLQPRPV*EML
						LL*PTLDLQPVVPPAVAILSWKP
						SALTISSQIVTPGLKTRSFFTRKT
						FMKIWNHRQKYQKTMLVMFP
				·		RYPSL
19319	49687	A	19432	599	900	ATWLFISPTTTTCQLSKTLLVPL
		l .				NFLEKQYLTFLLFLEFPTFSLFF
	}					RHTEDHLMRSGCLKVCSISPLS/
	1	ŀ				PLPPSSSHVRHLTPSLPFAMIVS
						FLRAPQKLSR
19320	49688	A -	19433	210		HFRKIFTVFQLKLVEYNLLSGH
1,7520		l' <b>`</b>	17.00	[-··	[	STQQSHPGGTVLL*DLVCPVV
		Ц		L	L	OT SACTOR AND DEACHAS

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19321	49689	A	19434	239	464	SPKGGPPNPSHGTKFHTRPCEE TTKQALCEEHGCLFHLGAGGLS PKRESAKGDKGGASVHVVCHI S*SPCS*LGP
19322	49690	В	19435	1	838	
19323	49691	C	19436	80	443	
19324	49692	Α	19437	1	775	
19325	49693	Α	19438	209	469	
19326	49694	A	19439	304	375	NVGE*SERHPCYD*TPREVCPPS P
19327	49695	Α	19440	879	1188	
19328	49696	Α	19441	615	917	ATWLFISPTTTTCQLSKTLLVPL NFLEKQYLTFLLFLEFPTFSLFF RHTEDHLMRSGCLKVCSISPLS/ PLPPSSSHVRHLTPSLPFAMIVS FLRAPQKLSR
19329	49697	A	19442	674	1554	GQMASSSVSLQQGVPQAADPG WVTAGASLTRAWAAGVQGDG PGTESSISASSRRVLRLGMTRTL LWACVLLWVCILLLGLCPAVG PVS\SVGPVSYCGPVPYCGPYPA VGPCPAVGPVPCCGPYPS/VGP VPCCGPVSCCGPVSCCGAVSCC GPVS\SVGPVSC/VLGLCPAVGP VS\S*CGP*PAVGLCPAVGPVPC CGPYPS/VGPVPCCGPVSCCGPV SCCWACVLLLGLCPAVGPVS\S VGPVPCCGPVPCCGPVSCCGPV S\VVGPVSCCGPVSCCGPV CDHSSHSLLLTFTML
19330	49698	Α	19443	4688	4824	

2426

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	l .	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19331	49699	A	19444	1	2274	MLKPLMGGGAHRRAAAGARK
19331	43033	^	17444	I	2274	WDGTAIIASTSGLEVLWVTTSR
1						SKTVESVGLAKPGQHNDARMN
						_
						LAIALTAARYGAATANYMEVV
					1	SLLKKTDPQTGKVRVSGARCK
						DVLTGQEFDVRAKCVINATGPF
						TDSVRKMDDKDAAAICQPSAG
						VHIVMPGYYSPESMGLLDPATS
						DGRVIFFLPWQKMTIAGTTDTP
		j				TDVTHHPIPSEEDINFILNEVRN
ŀ						YLSCDVEVRRGDVLAAWSGIR
						PLVTDPKSADTQSISRNHVVDIS
						ESGLITIAGGKWTTYRSMAEDT
						INAAVKTHNLKAGPSRTVGLFL
						QGGKDWSPTLYIRLVQDYGLE
						SEVAQHLAATYGDKAFEVAKM
						ASVTGKRWPIVGVRLVSEFPYI
						EAEVKYGIKEYAC/DCCGYDFT
]					ļ	SYSPGLS/EMSRQQRKPYPGLL
						N*/CGRELNWDDYKKQEQLET
						ARKFLYYEMGYKSRSEQLTDR
			:			SEISLLPSDIDRYKRRCHKFDAD
			ŀ			QRGFMTVVEVQRVLESINVQM
			•			DENTLHEILNEVDLNKNGQVEL
			:			NEFLQLMSAIQKGRVSGSRLAI
						LMKTAEENLDRRVPIPVDRSCG
						GLYFCVRNWWVLGLTDFKNE
						AADPSGVKLQTFTVSVTARKGS
						VDPKNSGAQLASPSGSRTGAA
						GGAACQSRALRLHSSALGWSM
						GLGAVEHGAVLIGEARAAQEP
						MEGVGGSGMAGCRSQVLPRG
19332	49700	В	19445	439	1542	
19333	49701	Α	19446	2	418	CLSLPPTPWAPVRPEPPRR/CAT
						PCSTAPSPIDHPRAEECERTVQD
						WRAAPPAAPVRDPLETGKIDQE
						IHKYNTPGFTGCLSRVQFNQIAP
						LKAALRQTNASAHVHIQGELV
						ESNCGASPLTLSPMSSATDPWH
				1		LDHLDS
19334	49702	A	19447	111	215	GTCWELE*RKGKPIFWGEIKAS
					·-	CRNLHKLQGAKC
19335	49703	Α	19448	740	959	DLYPHHLSWQSRNAARKRKS*
1,7555	1.7703	<b>'</b>		1, 10		RGQRPIPAIS*GR\SQKQQQNVN
						TSTPSHCPRLPYECRPLLHGAQS
						HRPPKG
19336	49704	В	19449	260	508	TAG TAG
19330	I	Г	17777	1200	200	<u> </u>

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19337	49705	A	19450	11	490	RLSTSPPDSGAQLASRSGSHTG AAGGAACQSRTMRSHSSALGW SMGLGAPLQRH*INSSGGSWLV DKESFPVPSALNFSPLGRKSSPC SKVLYMPNPVSHSMIPSGSERS MDTNGSCWYLTCSLSAPGRRD TRIRKVFSQGRLLVLTLMISPSV GHSTA
19338	49706	A	19451	42	263	SGVSPMSFNTFTSAPAFNRSWT TCLCPK*HARCRGVQPKLDLLL PLQLMSAITSFLAMVSNSCLTSS ALPFLA
19339	49707	Α	19452	662	1564	GSRICSFQTRSGHPECPQCHWT RQNHSSIQQPHKAAYTKKTTIH PPNRSNSCSSIWCKLQKLYMCG P*GRSFNFSANLHQPQKGHSGE RPF\EKNKGRAILLKSHRVHTQE MPFPCSKVGKGYLVSSGLFQH QAIHNEKPCRSAMYGDMFHTQ QGHFKCIDYGEAFSPKDTPGQH QIIHTGEKPYVCTECEKTCTRSS NLIQHKPKGLLSLLVLSPSTGN ALFSTLSQLTQLLLTFDVRSDA TFFRKGSGRHQLWLALPEDAR ARVTSVSGTRRRDHGGARGVV CLIHGCVPDTRSSPDVW
19340	49708	A	19453		1167	MIREQKVKEVYICAGKKKSHM HLWHNGEKYNRLGRKDRKMT HEGWLIKIPHFPGRKYKASGLH WEQMKHMGGRGKVHLEFTEM YVAGRRGEVKDEVLEPMSLTE VYIPTENDREVWRERRLRAPGL CAALEGQLEFRVGVGLAGSAL EAAGQPCWPRAGRARDLQPAM PEPPPASVGSCAARASLRSAAP CSTAPSPIGHPRAEECERTAQD WQAAPPAAPVRDPLGEASWAP ESVMWPPLDSRQSEEVSVSKRH MDATDKIKFPWLKEKRTIEFGE AAGYVCHRRYKDMEPHEKMG EAAKKVHWVQMRSRARDLQP AMPEPPTHSKGSCAARASPT/T HHPLLHDAQSHRPPKGLRSAGT WRGTGRQLHLQPQCGIHWVKP AGLLSLL
19341	49709	В	19454	118	559	

SEQ ID		1	SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence	1	09/540,417	sequence	of peptide sequence	detection, \-possible nucleotide insertion)
19342	49710	A	19455	26	813	LSHYLACSLRLPSGSRIRSAWLI
						PSIMLLNYTSPGLPRHRSKSENR
						LDDTARRSPGMAGPGCPPLPGG
						VPDVRAAYGAPTSAQRPRVGW
						PPGGRWPRGARGARSFLKKPFR
		1				GTPTQVRGAWQCPPPPLTV\EK
						*PWHPCGRFQGRRHEEKGGRE
						LGHSKSPPPRRSHYPAGEKGEE
						ERKEREESSCAGLRRGAHEHA
						QGQSLPGELPAPGDPHPGRCVG
		İ				GQGRSRMGPCS*SLAQHPPVCT
						RGQREKQERRATNKPVRGAPK
19343	49711	A	19456	497	756	TWLKSGISFPPALFTIAKTWR*P
						KCPSMIDWIKKMW/HIYTMEY
						YAAIEKDEFMYFAGTWMTLDT
				ļ		VILSKLSQEQKTKHRMFSLISGS
19344	49712	Α	19457	100	819	EWSSVRRILVEKRALRRPHPQC
					İ	LCFRMKTILSNQTVDHSQKNV
						DITLKGTPQ*SCKGPRGT\LR\RD
						FN\HIKCGTSALLGKEQQRGFR
						VD\KWWGNRKGNWATRSGLF
		İ				GSHVQDHDPRGVTTGASGYKD
						EGPVYASPSPHPTVVIPGRNGSS
						LLKSRNFLG\EKYIPQGFRMKTR
						VLLCSVSQSPEKMKLILE\GNDI\
İ						ELVFKFQRALIQQ\ATTVKNKG\ IRKFFGWVSMSLEKGTVQPGLI
19345	49713	С	19458	395	574	IRRITOW VSMSLEROT VQFGLI
19345	49713	A	19459	1	1476	
19347	49715	A	19460	2	323	
19347	49716	A	19461	539	711	HRPFPQDRKAGECLLHEYEDLV
17540	13710	``	17101		1	PIRDTLRLFPGGRYLPRAKHVA
1						PSEPDPGWLVLG*FYC*FCHSP
ł						ALLSGFCLAVNVYRERSV
19349	49717	A	19462	66	2873	SARTISYDYYQNWGRDGGPRSS
1.75.17						GGGYGGGPAGGHGGNRGSGR
						GGGGGGGGRG/WQGPASRAPE
						RPRNRHVVREKTGAEEQGSWR
						GKREL/LVHMDERREEQIVQVL
1						DSGQAENG*RVRTTDF\RFAPE
						DHGYGTEVSTKNTPCSENKL/G
						HPGKEVDKSRKKNV/SESGTDH
						ILTRDSEYLLQENEPDGTLDQK
						LLEDLQKKKNDLRYIEMQELV
						NLIDNHQVTVISGETGCGKTT\Q
						VTQFILDNYIERGKGSA
	1	<u></u>	l	<u> </u>		VIQFILDNYIEKGKGSA

SEQ ID NO:	SEQ ID NO: of peptide	Met	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19350	49718	Α	19463	1	2112	FRKTTRRLRKVKGRTAGCARF
						RKWLLRQHGLRTLGLICNHTSR
						MTRCRWSAESDSGDEEEEGPIV
						LGRRQKALGKNRSADFNPDFV
						FTEKEGTYDGSWALADVMSPL
						MKKRAATTLHEKIEKARKRRK
						TEDKEAKSGKLEKEKEAKEGSE
:						PKEQEDLQENDEEGSEDEASET
						DYSSADENILTKADTLKVKDRR
						EEEEERTGSRRIF*RCIS/ATMKT
						SSFQDMNLSRPLLKAITAMGFK
						QPTPIQKACIPVGLLGKDICA\C AAPGTGKTPPFALPVLERLIYIP
						SPGSSL/TRVLVLVPTRELGMQ
						VHSVTRQLAQFCNITTCLAVGG
						LDVKSQEAALRAAPDILIATPG
						RLIDHLHNCPSFHLSSIEVLILDE
						ADRMLDEYFEEQMKEIRMCSH
						HRQTMLFSATMTDEVKDLASV
						SLKNPVRIFVNSNTDVAPFLRQ
						EVIRNRPNREGDREAIVAALLT
						RTFTDHVMLFTQTKKQAHRMH
						ILLGL\LGVQVGELHGNLSQTQ
		-				RLEALRRFKDEQIDILVATDVA
						ARGLDIEGVKTVINFTMPNTIK
		ł				HYVHRVGRTARAGRAGRSVSL
						VGEDERKMLKEIVKAAKAPVK
		1				ARILPQDVILKFRDKIEKME\KD
						VYAVLQLEAEKEMQQSEAQI
						NTAKRLLEKGKEAVVQEPERS
						WFQTKEERKKEKIAKALQEFDL
10251	40710	ļ. —	10464	254	116	ALRGKKKRKKFMKDAKKKGE RQHAH*ESSPLPNLTYPGTQILR
19351	49719	Α	19464	354	446	KAAGTSA
19352	49720	A	19465	42	388	RAAUISA
19353	49721	A	19466	177	497	
19354	49722	A	19467	271	773	TAMSSEEGKLFVGGLNFNTDE
	"   "					QALEDHFSSFGPISEVVVVKDR
		1				ETQRSRGFGFITFTNPE\HASSC
		1				Q*RSQ*TGEVSWNG\RQIPLLDH
		1				AGKSCSGEPRGRWALGPMGVG
		1				SQLILEVVGDQG\YGSGRYYDS
						RPGGYGYGYGRSRDYNGRNQG
						\GYDRYSGGNYRDNYDN
19355	49723	Α	19468	2	414	GRVGPPRRAGGPRASAAGDPA
						RGGAAPKMSQTAMSETYDFLF
						KFLVIGNAGTGKSCLLHQFIEK
						K/WSVTRSYYRGAAGALLVYDI
						TSRETYNALTNWLTDARMLAS
						QNIVIILCGNKKDLDADREVTF
	1	<u> </u>	<u> </u>	<u></u>	<u> </u>	LEASRFAQEN

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
NO.	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19356	49724	A	19469	397	445	GHRGAHRPRTLRSVVVRRAHR CSYSGIWD/TAGPDRFRSVTRSY
		1				H*GRPGALLVYDITARETYNAL
						TNWLTDARMLASQNIVIILCGN
						KKDLDADREVTFLEASRFAQE
						NELMFLETSALTGENVEEAFVQ
ĺ						CARKILNKIESERMGSGIQYGD
ŀ						•
19357	49725	A	19470	2	559	AALRQLRSPRRAQAPNAQECG
19357	49726	A	19470	1	483	
19359	49727	A	19471	95	352	RKATKMATIWASGTPKQPCPR
17337	49121	^	19472		332	ELLAQERRQTRTRYNQ*DQ*QE
1						RRSRQRIMGVGCLHMLRSEKA
			ŀ			PQQRSFAHCTPKSSRPSARLT
19360	49728	В	19473	247	1364	I QUELLING TRESID SAILET
19361	49729	A	19474	422	877	CHTQTQ*QHTP*QPTPIDMAPTS
	' ' ' ' '					HARMPFHTHFLTQILTLVSRSP
						DPG/PHGGKVPTRIPIPPAQPWG
		İ				PSTLPSSSFLGPPRPFPSPPVPGP
						GGENRKPFTSPP*A*\GFPGPPSG
ļ						P*AGGEG/PPVVGRGATPSPPTP
						SPACGPLKCWWHLIKN
19362	49730	A	19475	168	522	LALHPTRPLIPATDDEV\TVTED
	1,3,23					KINAPH*KQPVVNVEPF\APGLF
						AKA\LANVNIGSLICNVGAGGP
						CCQPAGGCTQPGGSCPLHCWL
						LPA*GRRKWEAKKEGFRGSFD*
						LTWGFGSF
19363	49731	Α	19476	1	328	
19364	49732	Α	19477	1	1204	MASRLLLNNGAKMPILGLGTW
						KGSILRVGFGSSSREPPPYRGQP
						LMGGPPVRILGPSGRPKRHNAR
						GRRGKWASQTGGPRAQTGTWS
						RRQGQVTEA\VKVAIDVGYRHI
			1			DCAHVYQNENEVGVAIQEKLR
						EQVVKREE\LFIVSKLWCTYHE
			1			ERGLV\KG\ACQEDTSSDLKL\D
						LPGPLTLIHWPTGFKPGKEFFPL
						DESGNVGPSDTHISDTWA\AME
			1			KLVDEGLVKAIGISNFNHLQVE
			1			MILNKP\GLKYKP\AVNQI\ECHP
	1		1			YLTQEKLIQYCQSKGIVVTAYS
						PLGLS*TGPWPKP\EDPS\LLEDP
						RIQGGSQPKHNK\TTA\QVLIRG
						SPCKRNLVVIP\KSVTPERI\AEN
						FK\VF\DFELS\SQGYDPP*LSLPT
						RNWR\VWCPCWSCT\SH\KDYP
					-	LPMKEVLKA
19365	49733	С	19478	300	489	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19366	49734	A	19479	243	471	CDHLQVSRPRIPATVPVEWKRS EMDSKEEEKEKKEKKEKRRR *RRRGKEEEEERRRRKRRKK RGRKK*IKKRV
19367	49735	A	19480	]	365	MGLYKDLLDTRYLFYLVPGYG LTQIELEVIAKDSAKPVGTPRV GLKLEMGIYLSGHLDKFIFAMP FKIKDILSLKKEEEK/ERRRRR RRKEEEDEEEEEEEEERRR RKRRKKKRKKKID*KKEEEKE EEEGEEGEKKKKMKKKRKKK KKKKGGGGGKGGERRRGRKK
19368	49736	A	19481	96	508	PAGSLVCPNQCSSARLFSSFFCS SAVMSASPAVGKRGWAELTGS FLPKPPLKTLGSGREDR*GQVW WLTPVIPALWEAEVGGSLEGQE FETSLGNMVKPLLYWLGSNGW LWKLGRRLWQLLWLQDDWPT RGSHILVL
19369	49737	A	19482	3	1013	SRPSWGTHWNEYARRLTLAPS WPLP*GHPVSPSLRVSCL/PHYY ESVLEKKGPCDRDYKKFWAGL QGLTIYFYNSNRDFQHVEKLNL GAFEKLTDEIPWGSSRDPGTHF SLILRNQEIKFKVETLECREMW KGFILTVVELRVPTDLTL\LPGH LYMMSEVLAKEEARRALETPS CFLKVSRLEAQLLLE\RYPECGN LLLRPSGDGARQRVGHHAADA QRDA/PWFRHYKVEAGRGPKY VIRCGTAVLLHLPWTTVVNYF VSHTKKALVPFLLDEDYEKVL GYVEADKENGENVWVAPSAPG PGPSPLHKVPPSRCQPASSQDKL PPLPPLRNPGREIT

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	i	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19370	49738	Α	19483	63	1496	LISAMAC\ARPL\ISVY\SEKGESS
ŀ			:			GKNVTLP\AVFKAPIRPDIV\NFG
						SHPNLRKNNRQPVCWQGN*AG
						HQT\SAE\SWGYWPRAVGSKFP
						R\VRGGGTHRS\GQGCFLETMC
						R\GGRNVLHPTKTLGRRWH\RR
	:			ļ		SSTQPKKRIRPSGSAPGCPQPLP
						ALGHGLKGHLY*GKFPETSFW*
						VER*KLKG*QEEPREAVFAPLR
	•					KLKRPGNDIQKGSMASSAK*RA
						GQKAKLRKPVARIPAAGAPCII\
			:			YNEDNGYSFKAF\RKHPWGLL
		İ				MLNC*GKLEHF*KLAPLVGHV
						G\RFLHLGLESALREV*DDLY\G
						T\WR*SHFPLQGVNLPIFSPWHK
						\MI\NTDS*AESLKSPEIPREPLRA
						PR\KKIHSQSS*RRNPLEKL*EIM
			ļ			LKA*TPFAKAHAAGTPFFGQAR
						N\HKLRVDKAAAAAA/ANLQA
						NSDEKAAVAGKKPVVGKKGK
			:			KAAVGVKKQKKPLVGKKAAA
						TKKPAPEKKPAEKKP\TTEEKKP
						AA
19371	49739	A	19484	3	730	KRLPKMAEVQVLVL\DGRG\HL
			<u> </u>			LG\RLA\AIVAK\QVLLGRK\VV
	:			1		VV\RCEGIN\ISANF\YRN\RLKY
						LGFSPPSGMNTNPSRRPLTTSG
						APSRIF\WRT\VRGMLPPQNPSG
						GQGRLLDRLQGV*RASPPPYGQ
			•			EKSGMVVP\AALKVVR\LKPTR
			1			KFCLIWGRL\APEVG\WKYQAV
						TAPPGGERGKRKAKIHYRKKK
	:		}			QLMRL\RKQAREETWRKKIDK
						YTEVLKTHGLPGLSPIKTVNSS
	1					KKKKKKRAAA
19372	49740	Α	19485	1	660	PTRPRTRGRRTRKSTPMKGLGA
						RYREYKV\VGRCLPTPQNATRG
						PLLPACEIFAP\N\HVVAKVPAF
						W*LCILS*KKMKKFFQGKIVYL
						WGKVF*EGSPLRVERTFGDLGL
						RLLTPRERAPHNIVTGEY\RGP*
		ĺ				PTAGGCSTQLLTRDNGVARHR
						RPEAHFHFRFMEGWEGDRGP/C
						KIPPGGLVKASSNDSKIKVSRW
						PHRV\LAAFRNKARFTT\KRPNT
L	<u> </u>		<u> </u>	L	L	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19373	49741	A	19486	51	471	EAGRLPGNILLDSIMACGLVAS NLNLKPGECLRVRGEVAPDAK/ S*VLPARCVHGILSDRLVGQWG WSWPRWPHAHPLPPSFVLNLG KDSNNLCLHFNPRFNAHGDAN TIVCNSKDGGAWGTEQREAVF PFOPGSVAEVCI
19374	49742	Α	19487	59	527	ETQSWPCGLVA\SN\LNFNTLES CLRFPKPSWLPDA\KSFRC*NLG QRPPTTWCLHFQPSASTAH\GS ARHASLCNSQGTAGHEGGPTQ REAVFPFQ\PGKCCRGVAFTFRT RANLNRSSLPDGITEFQVSPNGF NL\EAINYMAA\DGGLSRSKCV AFD
19375	49743	Α	19488	2	440	KQQCRDTGDEVVCSCFVGYQL LSDGVSCEDVNECITGSHSRAF GESCINTVGSFRCQ/RISSCATG YELTEDNSCK\DTLGSFRCRPKL QCKSGFIQDALGNCIDINECLSI SAP\CPIGVT\CINTEGSYKCQKN VP\NCGPLVNHLKE
19376	49744	A _	19489	2	2061	
19377	49745	Α	19490	2	2120	
19378	49746	В	19491	63	255	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19379	49747	A	19492	2	1941	PPPMERAAPSRRVPLPLLLLGG
						LALLAAGVDADVLLEACCADG
ļ						HRMATHQKDCSLPYATESKEC
						RMVQEQCCHSQLEELHCATGIS
				1		LANEQDRCATPHGDNASL\EAT
						FVKRCCHCCLLGRAAQAQGQS
						CEYSLMVGYQCGQVFQACCVK
						SQ\ETGDLDVGGLQETDKIIEVE
						EEQ\EDPYLNDRCRGGGPCKQS\
						CRDTGDEVVCSCFVGYQLLSD
		1				GVSCEDVNECITGSHSCRLG\ES
						\CINTVG\SF\RCQRNSSCGTGYE
				i		LTEDNSCKDI\DECESGIHNCLP
İ						DFICQNTLGSFRCRPKLQCKSGF
						ITRML*GNLLLDIQWSVLSIQLP
						RCPIGD*HGI\NTEGFYNVPEGT
						CPNCGPLGYHLQRGRGTRCV\D
						V\DECAPPCLSPVGKG\HR\CVE
				•		LFPGSFPLAECKDGVTIFDGISR
						DVLFDVQRSCQRLPRGACC\GH
						K/CTENTLGSYL\CSCSVGF\RLS
						VDGRSCEDINECSSSPCKPGSVA
		ł				NV\YG\$\YQCY\CR\RG*PASAIV
				1		GWEFTCERTFDGVAALPHRGH
		1				TFWFLTAGIQTSWGSFPVAACP
						SSGLTGWAPNGPATCQ\DI\DEL
						C*LGIHNCFHSTEDLVSNIPGAG
						FRCLGLSSCPENYRRSAAT*S*G
						NSCMRAIGAGWEKEKGKLAGS
19380	49748	В	19493	123	237	
19381	49749	Α	19494	285	456	VINKRLVCTRLLRRKWSCLQPH
		1		1		RMM*VQAGSRGSQPKYHRHE
		$\bot$		<u> </u>		MGWTGHTDTPPHRL
19382	49750	A	19495	383	693	NFLPAQPCLSAPPS*S/RSLLDNS
		1				PLNCKAFHSSSHHLLKLESCSE
		1	]			NSPGPLL\QPGGTCLGACPGCFS
		1				FSPGISAGKPLSKAASPPVCTGS
						GAETSTTPPSPLR
19383	49751	В	19496	28	236	

SEQ ID NO:	SEQ ID NO: of peptide sequence	)	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19384	49752	A	19497		927	PAAPQVAGRLGLGCPLHLHVF AVVSAMLPLLRCVPRVLGSSV AGLRAAAPASPFRQLLQPAPRL CTRPFGLLSVRAGSERRPGLLRP RGPCACGSGCGSLHTDGDKAF VDFLSDEIKEERKIQKHKTLLK\ MSGGWELELNGTEAKLVRKVA GEKIPVPF\NINNSIPP\TFDGEEE TLARARRFEDREPELTSLPKF\V VE\VIKND*WQERPLVLD\VHYP EDEVGQEDEAESDIFSI\REVSFQ S\TGESEWKDTNYTLNPDSFWT GPLYD\HLMGFP*PNRGVDNTF
19385	49753	A	19498	3	342	A\DE\LVELSTAPRSPPGVPLPFP LDTNLPANRVPAGLEKRLCAA AASILGKPADRVNVTVRPGLA MALSGSTEPCAQLSISSIGVVGT ARDNRRH\GAPFFGFSPRELALG QDRILIRFFPLESWQNGKIRTGM TFL
19386	49754	A	19499	3	520	SSARAVSVPLPAMPFLELDTNL PANRSARGGWRKRLCAAAAS\I LGKPA\DRVNVDRYAGGFAIGA *AGST\EPCA\QLS\ISSIRRQWGT AED\NRSHSAHFF*VSSPRELSP GARDRIILFRFFPLESWADWAR *GTVIDFFMIWGTEGFQGHLVN LAASFQKRSSWAEVKGLG
19387	49755	Α	19500	7	355	-
19388	49756	В	19501	172	479	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	1 ··· • · · · · · · · · · · · · · · · ·
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19389	49757	Α	19502	1	1656	LEDPDSSSRPRRAGSGHSASFPF
				•		PPLEQLWRQVKTLNPKAEVAR
						AQAALAVNISAARGLQDVLRT
						NLGPKGTMKMLVSGAGDIKLT
	}					K\DGNVL\LHRNCKFQHPTASL\I
						AKVSNKPQE*YNLVDGTDFLIV
						LINWESLLKQA\DLY\ISEGLHPR
						IIT\EGF\EGAKEKALQFLEEVKV
						SRE\MDRETL\IDVARTSLRTKV
						HAELADVLTEAVVDSILAIKKQ
						DEP\IDLFMI/SVIMEMKHKSETD
						TSLIRGLVLDHGARHPDMKKR
ŀ						W\RDAYILTCNVSLEYEKTEVN
						SGFF\YKSAEEREKLVKAERKFI
						EDRVK\KIIELKRKVCGDSDKGF
						VVINQKGIDPFSLDALSKEGIVA
						LRRAKRRNMERLTLACGWG/M
		İ				PLNSFDDLSPDC\LGHAGLVYE
						YTLGE\EKFTFIEKCN\NPRSVTL
		:	:			LIKGPNKAHTSLRIQRWQ*GDG
						L*GAV\KNAIDDGCVVPG\AGA
						VGSGQWAGSPGFKHKPSC*RA
						RAQLGSPKHLLDAFAHYFPKVL
						CFRNSGFLTFQGNIKLKFKAE\H
						SESG\QLVGVDP*TPGEPNWWP
						ARSKAVWG
19390	49758	Α	19503	2	424	EGPVMAPPNPGPATLGGPGPDP
						DLGRLPLHEVFLHIRVPARPRG
						APLHRRGLRGRHAVRAVRQRR
						REPEDGAAGAVDRAGGAGVLG
						RGDRESEGPLTD*PREPADRAP
						LLQPERGRFSHPLPTLTCAPSPII
						FPVAERWG
19391	49759	Α	19504	3	174	
19392	49760	Α	19505	2	153	LPLHEVFLHIRVPARPRGAPLH
						RRGLRGRHAVH*PSEPGDPARL
						LQPERG
19393	49761	A	19506	3	354	EDGRHGAPNPPPATLGGPGPDP
1		ĺ				DLGGLPLHEVFLHIRVPARPRG
						APLHRRGLRGRHAVRAVRQRR
						REPKNGAAGAVDRAGGAGVL
						GPGDTEYEGPLTD*PSEPGDPA
						REPKNGAAGAVDRAGGAG

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19394	49762	A	19507	3	1246	GSTHASARMAVMAPRTLPPAT LGGPGPDPDLGGLPLHEGF\LHI RVPARPRGAPLHRRGLRGRHA VRAV\RQRRREPEDGAAGARG* EARKGPE*LGRG\NTESEGPTHR LDPSEPGDPARLLQP\E*GRFFTP FQIMYGC\DVGSDGAFSSAGYP QERLTNGKDYIALNE\DLRSWT AADMAAQITKRKWETPPMRRR SLRAYLDGTCVEW\LRRYLENG KETLQR\TDPP\KTHMTHHPIS*P *RPL*RCW\AL\AFYPCGDHT*P WQRDG\EDQTQGHGSLLETRPA GAG\TFQKWASMVVPSGQEQR\ YT\CHVQQ*RVLPKPLTLKM/W EPSSQAHPSPIRGASFAGPWFSL GSL*ITG\AVGRCW*C*RKEQLR WRRGER*KKEGAYSQGCKASD
10205	10762	ļ	10500	1	202	QCPGASDVSLHSFVKVLRQLAL
19395 19396	49763 49764	A A	19508	1	392 220	
19390	49765	A	19510	3	500	
19398	49766	В	19511	28	134	
19399	49767	A	19512		1662	MAASVVCRAATAGAQVLLRA RRSPALLRTPALRSTATFAQAL QFVPETQVSLL\DNGLRVASEQ SSQPTCTVGVWIDVGSRFET\EK NNG\AGYFLE\HLAFKGTKNRP GSALEERRWRAWGAHLNAYST REHTAYYIKALAKELPKA\VEL LGDIVQNCSLEDFTD*GRNVDV ILRE\MQEN\DAS\MRDVVCNYL HATAFQG\TPLAQALEGPSENV RKLSRADLTEYLSTH*QGPSKW CWAAAGREWEHQQLFRTFAQE GTSVGIP\WTYAEDAVPTLTPCR FTGSEI\RHRDDALPFAHVAIAV EGPGWASPDNVALQVANAI\IG H\YDCT\YGGGVHLSSPLASGA VANKLCQSFQTFSICYAETGLL GAHFVCDRMKIDDMMFVLQG QWMRLCTSATESD\VDRGKNIL RNALVSHLDGTTPVCEDIGRSL LTYGRRIPLAEWESRIAEVDAS VVREICSKYIYDQCPAVAGYGR SLGLPFHLCSRTAAQLTEQPVK APSSSPQSPGCQGVLAQKTCPSS SLTACQGAFINTLHPSGPIEQLP DYNRIRSGMFWLRF

SEQ ID			SEQ ID NO:	1		Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	l .	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19400	49768	Α	19513	2	1693	AQSTLARRNRADCSWKMAASV
						VCRAATAGAQVLLRARR\SPAL
						LRTPALRSTATFAQALQF\VPET
						QVS\LLDNG\LRVA\SEQSSQPTC
						TVG\VWIDVGSRFETEKN/DSGA
						GYFLEHLAFKGTKNPALAVPW
İ						EKEVESMGAHL\NA\YSTREHT
						AYYI\KALSKDLPKAVGALGDI
		ĺ				VQNCSLEDSQIEKERDVILREM
						QENDASMRDVVFNYLHATAFQ
						G\TPLAQAVEGPSENVRKLSTA
						DFTRYL\STHYKAPESVLPQTEG
			ļ			VRTEQLLDL\ARNHRVHIP\*TY
						AEGAL\PTITPCRFTGSEIRHP*D\
]						ALPFAH\VAI\AVEGPG\WPAPD
ŀ						N\VALQVGNA\IIGHYDCTFGG\
						GVHLS\SPLG\SGACGPTKLCQS
						FQTLS\ICYAEDGLCWVHTLFC
		1				DRMK\IDDMMFVLA/QGHWMR
						L\CTSA\TRSEV\ARGQKNIPQK
		l				MPLGISS*DGNYSLCVEDITPAA
1						LLTLWARRNPPGLKWES\RIAG
				1		GGFQVVVR*DLAPRYIYGPSCP
		}				AVA\GYWPH*SMLPDYNRIP*P
İ		1				ACFWVAFFRAGKPMVKQKGR
						GGGLWSPPPPQTQHFRLLLTCC
19401	49769	Α	19514	46	422	
19402	49770	Α	19515	568	1567	GPWAANGVVLAFPPCLGSPPAP
						HLAPQKAVPPKWGISSLCSFHF
1						HIYQVGGVAFCGRIRFLSSTDF
						KRRAGGSGSLMSQEEKHELSG
		ŀ				KIRSQIMDDHARPYLYTMSNCP
						CWAGALGPQESKCSRGRLLHR
	1	l				AFPSLV\TLLLAGQATTGLLSCY
			İ			QQQGAGWDKLDSHRPRNLAA
						WEEPCGMEAFPKPSQSLVEAKI
						AQLATPGLLDCKAACPWGAPC
		i				PRGP\MQNATKVIGNM\TEDHV
		1		}	1	DAPGSRNAEPP*KVYPP\LKGSF
						P/EKNLRHLKNTMETIEWKAFE
						NW\MHHW\LLF\ELSKQSLWEQ/
1						QSPLDAPPK/ENSLELEDPSSGL
	1.055	_	10.51	<u> </u>	1.400	GVTKQDLGPVPM
19403	49771	В	19516	1	1428	

SEQ ID NO:	SEQ ID NO: of peptide sequence	l .	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19404	49772	A	19517	1	2293	   MARSEAAAAGPGPGPPGLAAR
	ŀ					ERRGGRGCVRGCERASVRVWS
					1	VALRRARRAPARPYVARCRRV
ĺ						RRPARVAIGCGGVSSPAGVSGC
						ARVSPERPPLRLCVWALSRAHG
						GAVCRAGPGPDSLKRGLTSHPP
		ĺ				AFPLHHGSRTLLLGPLSPPPAAL
						PFLLLLWAGHLVAS/ACPGRCIC
						QNV/SPTLTMLCAKTGLLFVPP/
						SIDRRVVELRLTDNF\RRHFAN
						MTSLV/QLTLSRNTIGQVAAGA
l						F/VDLRALRALHLDSNRLAGC/
						GGDQLRGLGNLRHLI/VGNNQI
İ						RRVESAAFD/DLLSTVEDLDLS
i						YNNLEA/LPWEAVGQMVNLNT
		ŀ			ł	LT/LDHNLIDHIAEGTFVQLQSW
1						A\RAQAAHPLTVSFGGNPLHCN
		:				CE/LLWLRRLTREDDLETCASP
	ļ					N/NLTDRYFWSIPEEESCTRCQR
						FCG*ASLVAAELTSNSVLIRW/S
						TQRPVPGIRMYQVQYN/SSVDD
						SLVYRKYSLSTYMNPTR\MIPST
						SQTFLVNDLAAGRAYDLCVL\A
						VYDDGATALPATRVVGCVQFT
		l				TAGDPAPCRPLRAHFLGGTMII
		l				AIGGVIVASVLVFIVLLMIRYKV
						YGDGDSRRVKGSRSLPRVSHV
		1				CSQTNGAGTGAA/AGPGPAGPG
[						PLRGAARGGVP/WLPPPSPSRPR
						PWRPR/LASAEPEVVLGRSLGG
						SATSLCLLPSEETSGEESRAAVG
						PRRSRSGALEPPTSAPPTLALVP
19405	49773	В	19518	97	255	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19406	49774	A	19519	1321	1728	EHAHDRDLHHAHYVQATLVR ASLREAVVGSPTDAWLQQRTA RSSASLLLARNRPATSVWLPRA GFIAGTGGRRERRGSEGWEGG GAGGTGAGLSAHSQTSACEGS/ CGGSAGRGRCLLSAATPPLQPP RPAPPTAGLVQPRQVPTAPPRA SAGNEHHQGWSGRL*KSLAAA RVGEAPGKLPRPRDFRKQSNGA LARASLRVSAQPGTCSSSAAAF PPLGPAPLAAPARSCDESGPRQP DGRGGPVPGQQQRGA/ESEPCA AATRPWVNRRR/TSRREARTS VACT*CAWCRSRSCACSHSPWS SASSSSAAICSSSPRA*STSS*RT GGRLRRWRRWSWGPTDPYAP AATAPTPAHFASP/ACAPHYQR LGEANLSESIISLD/CRPFGKKRP VSPSPSESPHAWQSEESAQTLN SAESGKKRAIRAQNFG/RLLLT WNAGRPTVRRPFSTRRWAKLW GQVERAGRGETQRH*SPCDRKS DLLKATQQTHGVSQIRVRVRSP LFSLLPPCPRKGGANKIPLNCKR
10407	40775	_	10520	102	447	ARFKKGFVFVCLEIQMFE
19407 19408	49775	A	19520	192	591	
19408	49776	-	19521 19522	281	421	
19409	49777	A	19523	174	512	
				67		
19412	49779 49780	A	19524		1166	MGAKDGRLLSESSQGLSSCGAS LVPFCVSKSRLLMKTPVRLDKE TVSTVDSGAPTDLAQLPTVLKQ PCCSVMASGQFVNKLQEEVICP ICLDILQKPVTIDCGHNFCLKCI TQIGETSCGFFKCPLCKTSVRK NAIRFNSLLRNLVEKIQALQASE VQSKRKEATCPRHQEMFHYFC EDDGKFLCFVCRESKDHKSHN VSLIEEAAQNYQGQIQEQIQVL QQKEKETVQVKAQGVHRVDV FTDQVEHEKQRILTEFELLHQV LEEEKNFLLSRIYWLGHEGTEA GKHYVASTE/PTVERSQEAH*FP VCLLASYDEISGQGASSQDTKT FDVALSEELHAALTSAHPDLILS QDEIVTLNLASQGGSQKRGNPR RFYRRGFLEVEPLTGFVELHSS GSQCQALIESGSREDLPVCPNK
19413	49781	Α	19526	203	359	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19414	49782	A	19527	135	466	DPAGSSAASMSSAAGLCALTPP
						/GLLSLGLLLLPLVVAF\ASAEA\
ļ						EEDGDLQCLCVKTTLPGSVPGH
						IHQLGG*FKGRDPHCPTCPT**A
		l				TVEGMGRKILLGTWQAPLLQG
19415	49783	Α	19528	46	468	TGPTHASAAILVPRSLHKMPGE
						ATETVPATEQELPQPQAETGSG
		l				TESDS/GKSPASDTYIVFGEAKIE
				,		DLSQQAQLAAAEKFKVQGEAV
						SNIQENTQTPTVQEESEEEEVDE
						TGVEVKDIELVMSQANVSRAK
						AVRALKNNS
19416	49784	Α	19529	337	509	ISQQLLVVLAGINLRSWLPQKL
						CKYLKQDVLTALDLSVEVPEFC
1		l				PFVDHLVVALEEEGPAL*GHPA
1		1				SDTYIVFGEAKIEDLSQQAQLA
						AAEKFKVQGEAVSNIQENTQTP
		l				TVQEESEEEEVDETGVEVKDIE
		i				LVMSQANVSRAKAVRALKNNS
	_					NDIVNAIMELTM
19417	49785	Α	19530	91	1019	GELLALLGHRVPTRPWASGQK
ł						GVCGWCPGKSTLQGLRCGPTIR
						ACRDGSSCSVEGGRGVRLVLG
						TASQDPEKEMPGEATEPVPA\T
		1				EQGVAASPRAETGSGTESD\SD
						ESVPE\LEEQDSHPGQPHNKAQ
						AGRQQSEIDEDPVS\KAKQSRN*
						KEGPGRAMSQTGVFGQVTGVT
		1				KEFTI\RKSKNILLCHPPKPDVY
						QGAPASRYFTIVFGGSPRSEDLS
		1				PSKHNLASCLRNSKFQGEAVSN
						IQENTQ\TPTVQEES\EEEEV\DE
						T\GVEVKDIE\LVMSQANVS\RA
						KAVPSPEERTSNDIVNAIMVSV
19418	49786	Α	19531	1 .	261	MWGHFHSHPGTGQARAYTDTS
						GSPIPLIMHVNRGRHSEIPVLER
				1		NLLHFRMDAST*NLGMSSLME
						AILKNQFRHVSLAVSFAFSGEK
19419	49787	Α	19532	277	349	
19420	49788	Α	19533	792	1014	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19421	49789	A	19534	849	1752	GESRGWERRKCVRPLSSYATHL SVCDCHIGNVREGADPAGGAG *L*GARQQQTHTHKPAKATQR GEDREEKKRETKREERLGQQE NERDERGRREKEPRNRGTKRRE AEKGRRR/ESEGKREEKGEEER QERQQKRAREKRRGRGARRRG REKKKTQRPRTQAQRPPLKKIK DYTHHTTQTPTQQHHTHTTKH KQPHTNNTTVFLFFFLLAKEHR RAARKQHEPERGRGGGSKAGA KIKHTSIAAAADRRHDRPTQRD TRRKEGRRAQSGKHGRTAHKT EAGGGIPVHSTSRVQVILLLQPP
19422	49790	Α	19535	139	470	
19423	49791 49792	A	19536 19537	32	434	NRRKGA\GPVFQRAT*KH/RV*R AGLRAVDFA\ERHGYIKGIVKDI IHNPRR/GAPLAKVVFRDPYRFK KRTELFIAAEGIHTGQFVYCGK KAQLNIGNV/LP/VGTMPEGTIV CCLEEPRLCRRKRTSAEGLNKV CVYAT
19425	49793	Α	19538	10	443	
19426	49794	Α	19539	63	376	
19427	49795	A	19540	170	368	
19428	49796	A	19541	175	427	
19429	49797	Α	19542	1727	2262	
19431	49798 49799	A	19543 19544	695	596 1337	GFGTQGERRRQCRLRVTQPLG ASSATQKTPALPETPVKAEQSC TLLPSPRTPGHLQPRLLDPSPW AVLIRG\QRKGAGSVVRRATLR HGK/GAVRRLLRRWISPERHGY IKG\IVKDIIHRPRSRRAPPLRQR WFFRDS\YRF*ER\TELF\IAARG AFHTGPSFVYCRQERPSFNIGQ WCSPVGHHCLEGYNSCCLPWK EETLETCGKAGPGAFKGN
19432	49800	Ā	19545	1	813	
19433	49801	A	19546	227	381	
19434	49802	A	19547	135	740	PQPLPPPTSMARHVFLTGPPGV GKTTLIHKASEVLKSSGVPVDG FYTEEVRQGGRRIGFDVVTVVR HPGGLLSRVGLEPPP/RENVNAE LGSYVVDLTSFWSSWALTRLE GM*ADCQQWAQGQRVCVIDEE LGKMEALSVK\FSFKAVRQTLS YPRDI*SL/VHKSQVPKGKPLAL VEEIRNRKDVKVFNVTQ\ENRN HLLARYS

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19435	49803	С	19548	251	574	
19436	49804	Α	19549	2	450	
19437	49805	В	19550	721	3048	
19438	49806	A	19551	3	396	
19439	49807	A	19552	2	450	
19440	49808	A	19553	2	1152	GPPLHCTRAAGIRHEAPLGLAG LCRAADAGLCSAGWGFVKVV KNKAY\FKR\YQVKF*KTPERV KRDYYARKR\LVIQDKNKYNTP KY\RMIV\RVTNRDINLSRLLYA PV*RGDMDSLAQR*LHTELAKI LVVEGLALTNLLPAALFVPGPA CLAPQAFLQ*VLAMDKDLLEG PSWS*LGDEYNV\ESIDGQP\GA FTCYL\DADLARTTTGNKVFG\ APEGAVDGGLSNPHSSKRFLG/ YVYPLHSTQTDSPGFMDSEKQR NFNARSTSGKHIMGQ\NCLQDY IALL*WERKMKDAYKK\QFSQY IK\NSVT\PDMM\EEMYK\KAHA AIRE\NPVLWKRRPKERSLKRR GWNRSQNVPLLRKKDRVA\QR
19441	49809	A	19554	3	991	KASFLQSSRSGLLESLTPAIFP LTSRGVGSCAVMLLLVRAGAR AWLRPTGSQGLS\SLAEEAARA TENPEQVASEGLPEPVLRKVEL PVPTHR\RPVHAWIESLRGFEHE RVCLVDLHPDVFANAPRLDILH QDAMLQKNF*RISYAKTKTRAE VR\GGGGKPLAAERH\WAGPA WQHPLSALARRRCCPWPPGP\T SYYYMLPMKVRALGLKVALTV KLAQDDLHIMDSL*L\PT\GDPQ LP*PELAH\YRRWGDSVLLVDL THEEMPQSIR\EATSRLKTFNLI\ PAVGLNVHSMLKHQTLVLTLP TV\AFL\EDKLLWQDSRLQTPLY PFSL\PYSDFP\RPLPHATQGP\A ATPYHC
19442	49810	Α	19555	31	305	
19443	49811	A	19556	213	541	DGGPAAHTPGAARGELSPVGD SGGYFRPSHLRCWILNLCISVPL SSAKDLLHPSSEEE\RRNPMKN RLVQSPNSYFMDVKCPGEETV CCIGESTGPQQLENVVVLAVSY

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19444	49812	Α	19557	164	591	DINSRDRSGPRATRIDGSGEGD
						KMAAQLRRHQGRRGASSPRFA
						KD\LLHPSPEE\EKRKHKKKRL\
						VQSPNS\YFMGV\KCPG\CFKFT
						TVF*AMAQTVVLCVG/CAAPLF
						LCQPTGEKARLTEGCSFRRKHA
				1		LKATLSQDEWETILQ
19445	49813	A	19558	362	916	GGRCTMARTKADWPANRTGG
						KAPKEANWLTKSPLRKSAALY
						LEGSKKPHRYR\PGTVAPP*NLD
		1				GYQKST\ELLIRKLPFQ/RRLVRE
		ł				IAQGLLKQDSRFQ\SAAIGCFAG
		ŀ				RQVRAYLVWPFLKDTNLCA\IH
			:			AK\RVTIYAKKTFQLSTPAYRG
į		ŀ				EPCFKDPLMDGETFHSQKKKK
						KISLLPVIGSSER
19446	49814	A	19559	849	1261	DGHIAGGPCS*GWPRDPAPKSQ
						PTMSSGSEPEKLPPAPRPQHYCS
ł						P\AEVSGPRFAVGIKALNQQAFT
				ļ		ERRLHTSLCPVPAALTVPSPESL
						PQPQIFPGLAPATHTLYSKAWA
						GATVSRPGAILGAGLYWGLVR
				İ		ETAS
19447	49815	Α	19560	3	1852	
19448	49816	Α	19561	362	1298	QGGRNVRHPSPLLAWGSDLIV
						KPHWLHWLLDLNHARLSALFA
						AAGARLPSHIPAPSAGSRRLLLL
						LLGGCRACLLFFPEMANRGPSY
						GLSREVQE\KIEQKYDADLENK
						LVD\WIILQCAEDIEHPPPGRAH
						FQKWLM\DGRVLC\KL\INRLYP
						PGTRSPYPRFSESKDGFLGRWE
						QISQF\LKA\AETYGV\RTTDIFS
				}		RRWFLWEGKDMAACAEGP*L
						ALGSVAVTKDDGCYRGEPSWF
						HRKAQQNRRGFSEEQLRQGQN
						VIGLQMGSNKGASPGGP*TGYR
						ECPRQIHVRTRHPAPWLERTNV
						STPWSFTKKK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19449	49817	Α	19562	3	2518	AAGEWPQRPRAAVAAAGAAA
						RORLPGSROVAAGAPOPAGGG
						DPSQR\PPGAEILIKVQVYVSGE
						LVPLARASVDVFGNRTLLAAG
						TTDSEGVATLPLSYRLGTWVLV
						TAARPGFLTNSVPWRVDKLPL
						YASVSLYLLPERPATLILYEDLV
						HILLGSPGARSQPLVQFQRRAA
						RLPVSSTYSQLWASLTPASTQQ
					·	EMRAFPCLPGH*G\SSSGNGSW
						LELMPLTAVSVHLLTGNGTEVP
						LSGPIHLSLPVPSETRALTVGTSI
						PAWRFDPKSGLWVRNGTGVIR
	]					KEGRQLYWTFVSPQLGYWVA
						AMASPTAGLVTITSGIQDIGTYH
	•					TIFLLTILAALALLVLILLCLLIY
						YCRRCLKPRQQHRKLQLSGPS
						DGNKRDQATSMSQLHLICGGP
]						LEPAPSGDPEAPPPGPLHSAFSS
						SRDLASSRDDFFRTKPRSASRP\
						AAEPSGARGGESAGLKGARSA
						EGPGGLEPGLEEHRRGPSGAAA
						FLHEPPSPPPPFDHYLGHKGAA
						EGKTPDFLLSQSVDQLARPPSL
						GQAGQLIFCGSIDHLKDNVYRN
						VMPTLVIPAHYVRLGGEAGAA
						GVGDEPAPPEGTAPGPARAFPQ
						PDPQRPQMPGHSGPGGEGGG
						GGEGWGAGRAAPVSGSVTIPV
						LFNESTMAQLNGELQALTEKK
						LLELGVKPHPRAWFVSLDGRS
		<u> </u>				NSQVRHSYIDLQAGGGARSTD
19450	49818	Α	19563	206	1126	LESGSHIYAFSHESLLLHVASAS
						V*TSFSDFATSVCHSPLHCQLH
						HALIPHGKGECSS\VSGIVATGF
						GATGFLGRCVVNHLGRVGSQIP
						QAIAQVSKEAEVGKFIHVAHLN
						VNIKSSSRYLRKKAVGEKAVR
						DTFPEAIIVKLLDIFGREDRFLN
	:					YFANMCWFGAIPLVSLGWKAV
						KQPVYVVDVSKGIVNAVKDPD
						ANGKTFAFVGPNRYLLFDLVK
						YIFAVAHSKGWVARDFEISPFE
						SWTTRDKVERIHITDMTLSHLP
						SLEDLGIQATPLELKAIEVLRLH
						HTYCWLSAEIEDVKLAKTVNI
Ь	L		·		1	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19451	49819	A	19564	196	553	KLSSEFFLTHSINTAIRNGNLFV
						NRVTEVTRSGGNELGSFCLLAI
						KFTLLSFLGTGTRRLLPALRPRR
	1	ĺ				LQQRGSHGLAGRLFTLWILHA
	-					GCHARGELLEPDTC*LSLCRAR
	ļ					KPRLSTLK
19452	49820	Α	19565	1	505	ATEDPGVAMGRRPARCYRYCK
						NKPYPKSRFCRGVPDAKIRIFDL
						GRKKAKVDEFPLCGHMVSDEY
						EQLSSEALEAARICANKYMVKS
	'					CGKDGFHIRVRLHPFHVIRINK
						MLSCAGAD/RIHISKKWGFTKF
						NADEFEDMVAEKRLIPDGCGV
				İ		KYIPSRGPLDKWRALHS
19453	49821	Α	19567	6	904	AQLSSLAVPLLVLTPFRSVLVFR
						SDSLFFVAATEDPGVAMGRRP
		İ				A\RCYRYCKTKP\YPKSRF\CRG
ĺ						CPLMPRIRIFD\LGRKKGKKWN
				1		EFPLLCP/HMVSDEYEQLSSESP
						GRLPRICPNK\YMVKSCGKDGF/
ŀ						HISRVRLHP\FHVIRIHKMLSCA
						GADKAPKQACRGAFWKSPKGN
						LLARGSQLAQVIMSIR\TK\LQN
						KEHVI*GPAARPSFKFPGRQK\I
İ				1		HI\SKKWGF\TKFQC*WNLKDM
						VAEKRVHPHMACGGQSTSPSR
						GPL\DK\WRALPLHEGLPNVAA
						HPLNTPPIKFLLPWSKKKKKK
19454	49822	C	19568	346	471	
19455	49823	Α	19569	1495	2780	EGLHLFCDLLDGLPGVQVAGG
						FDSFSSILASFTY WHDFPNSTML
						LSTFRDINLGMNIIKKSAVCFSR
				}		LGSETWEMPANLRPSEGGART
				1		RTELSRPQTPSRAHLAWRGRGG
						LGAVRPRRLSRRRSLS*KP\RSL*
						AEAPAGAPVRRCRPAAPP/PPRP
						SRCGPSTPPPPPSPGA\PRRRPP
				1		CSEHNKLSPPPPSSLCLRPAPSPP
				1		RPAPRRSPSRRRRRRRHPRPAD
						RREPLASATRVRPLPPPPPPR*PP
						SSSCPRRGGRRFLAPWLARPAG
						APRRAPCSRAARSRRAGASGAR
				1	:	VGDPAGAGRGWRRRAAGRP\*
						APERRPPPPLPAVCEVSGGPLRA
						SFADSVPGVRRRRPPGLGTPLSS
				1		SGV*PRNERGVETTTSWGGGSE
						GAPGSGEGGNLFPAPQVIVARS
						GEGTVVPRFPVCQLRDSAVLLC
						LFD
19456	49824	A	19570	178	256	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19457	49825	Α	19571	622	1726	QGSGRTMMRMLVPASVGLLET
						GPPGDPSLSHPLSGDVTVLPED
						Q\STVQKLASDHKDIHSSVSRV
						GKAIDRNFDSEICGVVSDAVW
1						DAREQQQQILQMAIV\EHLYQQ
						GMLSVAEELCQESTLNVDLDF
<u> </u>						KQPFLELNRILEALHEQDLGPA
						LEWAVSHRQRLLELNSSLEFKL
						HRLHFIRLLAGGPAKQLEALSY
						A\RHFQPFARLHQREIQVMMGS
						LV\YLRLGLEKSPYCHLLDSSH
						WAEICETFTRDACSLLGLSVESP
						LSVSFASGCVALPVLMNIKAVI
1						EQRQCTGVWNHKDELPIEIEL\S
						*SAWYHSVFACPILRQQTSDSN
						PPIKLICGHVISRDALNKLINGG
19458	49826	В	19573	271	392	KLKCRYCPMEQNPADGKTH
19459	49827	В	19574	58	296	
19460	49828	A	19575	215	861	QLPAESVRTRGPAKAQARAGG
17400	17020	<b> </b>	17575			PPGPWHCSYKWRAWPSSLSGT
						GWPDMEGETTAAAPASRENSA
						DCS\AGELRGPG\KELW\SPIQ\Q
						VTATSS*MGRDFVLAT*EKVSH
j						VDS*/EQPRSLQR\DPRP\AGPAQ
						AKQGTPRAQASRESL\SRPTAA
						V\QLPSGPQHPV\TSGS*KGLAG
						KTSMGRKGLPWARGWAPWSL
•				ĺ		EAQNP\RPTRLCDLFITGEDFDD
						DV
19461	49829	A	19576	65	782	SSSSSAIQRKPRWVLYKYIQEL
15401	77027		17570	05	7.02	WRKKQSDVMRFLLRVRCWQ\Y
	İ	İ				RQL\SALHRAPRPTR\PDK\ARRL
						GYKAKQGYVIYRIRVRRGG\RK
				ŀ		RPSFLKGAT\YGQSPVHSWVFN
ŀ				ŀ		QLKSFASE/VLSVPFARGRRSLG
	1					RH\CGGLLEEFLNFFNWVWVER
				ŀ		FPHYQIFLKVIPHLIPFP*KLFRR
		1		}		N\P\DTQLIHQNQVHKHR\EMRG
1		1				\LTSARPEKSRGLGKGHKFHHTI
l		1				GG/SLRRATWRRRNTLQ\LHRY
						R R R R R R R R R R R R R R R R R R R
10462	49830	A	19577	1	179	QEDYLRELLTTMGDR\FTDEEV
19462	49000	A	19311	1	1/3	DELYREAPIDQKGG/MFNYIEFT
		1				RILKHGAKDKDGLKEL
L	<u> </u>		L	<u> </u>	<u> </u>	MILMIUMNDALKEL

SEQ ID		Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19463	49831	A	19578	144	722	KKGLNHHHVEQNNTRPRPRSG
						PQRATSN\VFAMFD\QSQIQEFK
	1	1				EAFNMIDPDQDGFHRPGKICND
						MLASLG\KNPT*WSILDAMMNE
		ł				\APGPPSNFTPWFPHHGFGEEV*
						MATRS*KMVIRKCPLTCFE*KK
		ŀ				PLGTI\GEDYL\RELLTTHGGIGF
						TDEEVDELYREAPI**KRGISITS
						EFTRILK\HGAKDKDD
19464	49832	Α	19579	3	429	VACLREHKQRCERATRSLLREL
						LQVRARVQLQGSELRQLQQEA
		İ				RPAAQAPEKEAPEFSGLQNQM
						QALDKRLVEVREALTRLRRRQ
E						VQQEAERRGAEQE/DRPQTGQA
						DRLAAAGGAGPGGGLRRPAEE
ļ		1				PRGQQPEGGPGGGQN
19465	49833	Α	19580	1	2454	RFPAGVAPRRAMANVSKKVSW
						SGRDRDDEEAAPLLRRTARPGG
		l				GTPLLNGAGPGAARQSPRSALF
						RVGH\MSSVKLDDELLEP\DMD
						PPHPFPKEIPHNEKLLSLKYESL
						DYDNSENQLFLEEERRINHTAF
						RTVEIKRWVICALIGILTGLVAC
						FIDIVVENLAGLKYRVIKGNIDK
						FTEKGGLSFSLLLWATLNAAFV
						LVGSVIVAFIEPVAAGSGIPQIK
						CFLNGVKIPHVVRLKTLVIKVS
		ļ				GVILSVVGGLAVGKEGPMIHSG
						SVIAAGISQGRSSSLKRDFKIFE
						YLRRDTEKRDFVSAGAAAGVS
						AAFGAPVGGVLFSLEEGASFW
						NQFLTWRIFFASMISTFTLNFVL
						SIYHGNMWDLSSPGLINFGRFD
						SEKMAYTIHEIPVFIAMGVVGG
						VLGAVFNALNYWLTMFRIRYI
						HRPCLQVIEAVLVAAVTATVAF
						VLIYSSRDCQPLQGGSMSYPLQ
						LFCADGEYNSMAAAFFNTPEKS
						VVSLFHDPPGSYNPLTLGLFTL
						VYFFLACWTYGLTVSAGVFIPS
						LLIGAAWGRLFGISLSYLTGAAI
						WADPGKYALMGAAAQLGGIV
		1				RMTLSLTVIMMEATSNVTYGFP
						IMLVLMTAKIVGDVFIEGLYDM
		1				HIQLQSVPFLHWEAPVTSHSLT
						AREVMSTPVTCLRRREKVGVIV
						DVLSDTASNHNGFPVVEHADD
						TQPARLQGLILRSQLIVLLKHK
19466	49834	В	19581	508	4413	

SEQ ID	SEQ ID NO:		_		1	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
		<u> </u>				
19467	49835	Α	19582	8	331	EVTACWQSSQPSLTLSASSAWA
	1					STLAALEEPFSPPLHSESPFLGW
						PRPELGPSAAPCAPVPPEPPQRV
						PPPAP*RPVPSTTQGLRSVGAW
		<u></u>				CRTGRQLHLHPGVGSTG
19468	49836	Α	19583	831	1121	
19469	49837	С	19584	1	1524	
19470	49838	Α	19585	270	578	RFSKSPPDSGAQLAAPSGSRTG
						AAGGAACQSRAVRPHSSALGW
						SMGLGAMEQGAVLVGEAWAT
ŀ						QQPTVGGRLRHGGLQVPSPAPR
10471	10020	<u> </u>	10506		200	GGS*GPAKSHSSCWPRC
19471	49839	A	19586	2	298	
19472	49840	В	19587	113	515	MI DONI DONI A MUDOD A CONTI
19473	49841	Α	19588	1	569	MLPDRLRSPVAHHRCRASGNY
						HSGRHSPAPAWRVTLTGWLDQ
						RSSRIHRSLAFKDSYSKGKEERT
	,					HIKGMCSELLPSG/EARGLAGFR
						SEAADLRGECYSS*RQCGPKEL
						RSPAGFTQWIPHWGYRWSCLS
						VPRPAPALLSPWVVDGTGRCG
						AGGGAHRGGSGRSGAHGGGG
10474	10040	<u> </u>	10500	1	1200	RLRHGGLQVPSPAPREGS
19474	49842	C A	19589 19590	511	1200 <b>846</b>	ARHRVLIGVFTIPELDIKVLHVP
19473	49043	^	19390	311	1040	TRLRSPASFTQWIPHWGCRWSC
						LPVPRRVPALLSPWVVDGTGG
						CGAGGGAHRGGWGCTGAHGG
		ŀ				GGRLRHGGLQVASPAPREGS*G
						PARN
19476	49844	A	19591	9	481	QQPLREKPASGDHPAKNANRR
17470		` `	17371	ľ	101	WPSPTRYPYVES*NWMPSLHLI
	1					QKSIQDGLKT*TLDLKP*KL*KK
						T*ALPFRT*A*ARTSCLKHQKQ
						WQQKTKLTNGI*LN*RASA/PA
						KETKI/SSEQATYKMGENFHNL
						LF*QRANIQNLQ*TQRDLQEKN
						KQPHQK
19477	49845	A	19592	1	3339	MSVRKDVEKLEPSDIVCGNVQ
						CYSCMETNLTVSQKVKHEVTV
						GPREGATKPNRMKGKEGRSGS
						LLGEGDFFKDESVMSSQGSSKD
						GEKRRGKAQRWKWPMQGICR
1						QLGVAKSMEGYQSRRDQGGR
						GVSDKWPQVCAKKPEFYPTAQ
1			}			VWANFSVTSCQSVTITQLCHGL
			1			RRLEISPARSNAMHLNPDPPGQ
						KQNLSPKVNDIITDIESSSGSGA
						GKFQVISKSDISEVLLQQMDAG
						HSSKDDPNEYGGWKSPRPRC
19478	49846	В	19593	1	631	
194/8	49840	ĪΒ	19393	11	031	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19479	49847	Α	19594	71	208	PCPKHCSKFPMCPELVGSWSH*
	<b>;</b>					LQE*SHGPSRFRLC*KQPQDLQ
19480	49848	Α	19595	138	650	AWAQLPSALGSTRVSPPPSARS
						QPPWRRRHKVCWRLNRCKE/L
						DEDAVASADFSSMLSEEEKEEL
						KAELVQLEDEITTLRQVLSAER
		l				KAS/ELR*NKNSA*T*/CNELKPE
						LQQKLA*HAD*HCLQEDT*YAE
						SRRAKGNCSFQQRWNGHQQEV
						RRHELLHSPFHKYACYETKVG
19481	49849	Α	19596	1	1222	MRADGTAGKDPFATDKQLPEL
						LIQHCWDGLLAVCGGSTVGVR
						LPFILHGSWVRPVAASFPPLPDN
		i	]			LHDSAEPAIILLEPTQMRRNQK
						TNAGTMTKQISSTLPKNHHTSS
		ŀ				PAMDPNKEEIHDLLEKEFSRLVI
						ELIREGPEKGKAQCEEIQKKDT
						KVKGEIFKEVDSLKKKQLNIQE
						NLETLLEVLNALESLSNRIEQVE
			:			ERNSEREDKVFELTQSNKDKA
						QRIRKYEPSLQEVWDYVKLPKL
						RIISFPNEEENSKSLENIFRGRIK
						EHFHSIARDLDIQIQEEQTTPGK
	:					LIVKRSLPRHIVIMRAPTPMDHP
		ļ				RAEECGRRARDWQAAPPAAPV
						RDPLGEASWAPDSEGRK\WNTS
						KHQRNKLRTRHL*EL*HSSRGS
						VASFLKSVRPRTHQFQTQSGYP
						KLRWRKES
19482	49850	À	19597	381	598	
19483	49851	Α	19598	1	1906	
19484	49852	Α	19599	493	589	
19485	49853	Α	19600	60	261	
19486	49854	Α	19601	1423	1831	PRRSANVNLPPPQLATLSPGWP
	1	l				EERKSVGSPTSPPPPRPLRRLSA
						APG\SGGAWPEPGAWSLD*TAS
						PAGAPVQRPAGNG/PSALDASH
						ARALYLQRIPRVTWTPASQPPA
						PLGPFLPLIPPAASHRQPSSSFFR
						RHR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19487	49855	A	19602	1	1535	MEEKTNVQLPPGQTEQHVEIHI
						MNFCSKNHHRITPEKPKELTDP
						FKEAACCCKLYEIDKKLYRMA
						EWIKIHKPSICCLQETHLTHKDS
						HKLKVSITFKDLAVRFSEEEWR
						LLEEGQREFYRDVMRENYETL
						VSVEPGRAVGGGSHADEGQEP
						AGCG/VSPGPGAAGEGDPRVLV
						WRSQGRYGQPRER\GRGASLD
				•		GERASPEAA/DGKRALPSPRPA
						QLPSRRPYQPAPPG\PTPTDSSCS
						SGPTGDGVQGSPLPIRISPGNSP
						L/PRPHQLSEGNPCAWAPAPRDI
						PKLLATSP*PGHVQANQSRPGA
						WEPALGRSDQRACSASGSAELC
						ERWPQQAP/APPEEPPPASPHPA
						APTG\PGFWESCGEPGAA\PGKG
				+		SAPKPSPLHCLESALRGILP\EGP
						CASPAWEAPAPAPAPAPARASA
						A/AEGEDPRPEPELWKPLPQER
						DRLPSCKPPVPLSPCPGGTPAGS
						SGGSPGE\APGEQSPGTAAASV
						Q/VSPAHWPCFS/SPVRYSSGSLP
						GFSAGEKAQG
19488	49856	С	19603	4	150	
19489	49857	Α	19604	148	448	RTRDWSLFLLMFGCVRSFFLLV
						GLWSCWLRSEAADLRGECYSS
						*GGASGVVRSSWWVRGLTSFR
						SETGFTQWIPHWGCRWSCLPVP
						RCAPALLSPWVVNG
19490	49858	Α	19605	136	439	LVHFTEC*LVRLQTLARHSTDW
						CIYKSLARQKSSPSPHSTQEVQL
						ASPLTGRQEDCSEPRSAPLHSSL
						GPTERDSICVAASSVRWTRPGA
						RFS
19491	49859	Α	19606	2	399	SKLQKLKRSLSFKTKSLRSKSA
						DNFFQRTNSEDMKLQAHMVAE
						ISPSSSPLPAPGSLTSTPARLVCI
						QVARLMPFRNTSSRSPLFC/RCL
1						QPHDSGFLFLRPARPRAHREER
						NNSRRVALRAVTLTTERLQLHS

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19492	49860	A	19607		1262	MPLSYAYKNAETLAGRHTSSW MSRGAYQRRNTRAAGRPEECT DRNWHAGRTRRIWLGQLEERC SDVFGVSFFWWVRGLAGSGAK LQTFTVSVTAHRGSKDQKSEQ KQDLLQKAKEQTIHNVKGDPS ELPLLAPAACFYSLICPRPPPHIL LIGPFYRELIGPFYRELIGRQLRP GEKSSTAA/VWPRC*AA/RLPGA G/WTAPSAGPAEPTPTWNSRWP ASATRRPGSRRRLSLHTSPQAE GAGSGLSQPRKGLPQCSGRLKR SSSAARVGVARQRRRERASEG/ WPACCHLSVLPCGFGFLEAHSS SPEQTDHEQNCPADIKVMRLA GQPTNARGAANRAQPLHSTSV SHLKAVIFKIQTRQQKNSHFPPL FLEMTTSLATGRLRPSWVDYA RPALSVVSVSRFSLLTSLLELLV ETLGECGLL
19493	49861	C	19608	173	394	
19494	49862	Α	19609	1	924	
19495	49863	A	19610	3	689	EVQEELEGPVRGAPHQLHQGP VEHLEAPHQ\HKFGKKVIYFNY LSELHEHLKYDQLVIPPEVLR\T MRSSRACTRAGRRLPPRHHRR GPRCPHSSLASVCNTSKTKIKA NSSPLC*GSQ*RT*ERKACAPRA CSGDPPACRPSARSRGSTT/KGK PVNFDDYGDIHIPAVILKTFLRE LPQPLLTFQAYEQILGITC/GGE QPACHSLPPDLTEPPRAQLRRPP LPHGLPACG
19496	49864	A	19611	1749	2411	APSCLVSEHSAPGPQRELPQPLL TFQAYEQILGITC\GSCP\AQGW GAWSSDAVPQLLARRPPLPHGL PACGEWGRGELGVKPSGLPSH AGPAWGHQVRTVCATAHPQD CISPEGAVEEEIVGG*GC\TEGQ SQRVLQIWP\SQGVSSLSALVPL N\MF\TELLIEYYEKIFSTP\EAPG EHGLAPWEQGSRAAPLQEAVP RTQATGLTKPTLPPSPLMAARR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19497	49865	A	19612	883	2164	EHRRPMRAQASHEGHRHPMRA
17477	17003	1	,,,,,,			QMSHEGYRRPMRVQASHENTG
						VP*GVQASHEGTGIP*EHRCPM
						RTQVSHEGTGVP*ECRRPMRTQ
						\GPMRGTGVP*EHRCPMRGTGV
						P*EAQASHEGHRRPMRGTGIP*
į						GAQASHEGHRCPMRGTGVP*G
						HRRP/DENTGVP*GHRRPMRAQ
						ASH/REHRCP/DEGHRRSP*GHR
						RPMRAQASHENTGVP*GAQAS
		:				HENTGVP*GHRRPMRGTGVP*E
						l .
						\QGVP*GHRRPMRAQVSHEGHR
ŀ						RPMRTQASHEGHRRPMRTQAS HEGTGVP*EHRRPMRGTGVP*E
						l .
	:	]				FRRPMRGTGIP*GHRRPMRAQA SHENTGVP*GNR/PYHEGTGVP*
		1				
		l				ECRRPMRTQASHEGTGVP*EHR
						CPMRAQASHEGHRHPMSTQAS
						HEGHRRPMRTQASHEGHRHPM
						RTQASHEGT\ASHEGHRRPMRT
		1				QVSHEGTGVP*EFRRPM/PGHG
ŀ						HPMRTQASHEGHRRPMRVQAS HENTGVP*GHRHPMRGTGIP*G
						HRCPMRGTGVP*GYRHPMRTQ
		j				ASHEGYR/PSHEGTGIP*EHRCP
		Ì				MRTQVSHEGTGVP*ECRRPMR
İ						TQVSHEGTGVP*GHRRPMRTQ
						VSHERHRRPMRTQASHEGTGV
						P*EHRCPM/MGHRRPMRTQASH EGTGVP*EHRRPMRGTGVP*EH
						RRPMRAQESHENTGVP*GAQA
						SHENTGVP*GHRRPMRAQASH
10.100	10055	<u> </u>	10610			
19498	49866	Α	19613		552	VPQDSLGQALAGQATPEIPLGL
						QLHTVLVQEIQELIEAQTRAPGP
						CAEVRALPAPAAEPEPAWEEAP
						PERALELEGAPAKDQTNEELPEI
j						TA/PYC/EPLALTLELKAWLERE
						VGGRG\DQHSPSQQLPCCP*SW
						ARWQTCRQRAGHLAWPPVPRC
						REASLIH*NHSPAAAGPFILLILV
10.100	1006	ļ. —	10514	1.12	001	YLCAGGA
19499	49867	Α	19614	443	991	VDSRPMLNTRLMRTSGRRWTP
						RPQVDTRP*VNNKTQ*AIRPQ/C
						GYSPQVPQDSLGQALAGQATP
						EIPLGLQLHTVLVQEIQE/RH*G
	:					TDTCSWSMCRSKSTASTSRWS
			}			Q/RPAWEEAPPER/GTGAGGSSS
						QGPDQRGAA*NHGGAGEH*KK
						AGRKSSE\RYACSQRQPPPPGM
		_			1.50	DAGPQLGIQAVCP
19500	49868	Α	19615	16	450	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hođ	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
İ						
19501	49869	Α	19616	2	477	PRVRSGHSMDMRVPAQLLGLL
ŀ				İ		LLWLRGANCDIQVTQSPDSLSA
ļ						SVGDKVTITCRASQGISNGLSW
						YQQKPGQAPTLLIYAASSLQSG
1						VPSRFSGSGSGTDFTLTISSLQPE
		1				DVATYYCLQDYTTPFTVLQAIT
1				j		KTPQR\RDV*VCTVLATPPGVIN
		l				WL
19502	49870	Α	19617	2	675	LPLSSSALPTESGSHLPFSGIMA
						RPSDPLVKPQRSSKREPKKVHP
]						GTQSDR\YVKIKR\NWAENPRG\
İ						IDN\RVR\RRF\KGQDLDAPTLG
ŀ		1				YG\SNKKTKAHACPSGFREVPG
ŀ		1				VHNVQGSWEVACLMLQQNLY
						CCRGSVHNVFLPRNRQSPSWG
						KELAQLGLSGVHPTPLARAAQL
						KEN*VRQLMCNKSYCAEIAHN
						VSSKNRKAIVERAAQLAIRVTN
		ŀ			:	PNAR\RSEENE
19503	49871	Α	19618	2	527	VWAANMPSRLRKT\RKLT\GHG
						AATGHGPHRPSTRKAPPGGPR*
						MPGGPAIHPPGSNFRQNTHPQA
						YFGK\VG**KHYPT*KRN\QSFL
					1	PPTVNL\DKLWTLVQ*NRHRVK
			]		İ	CLPKNKTGAAPHHLMLVRSGY
					•	LQSSGERGKLPKASLFFVEGPK
						FFSRRSLREEGLRGVWGGPVSL
19504	49872	Α	19619	45	233	
19505	49873	С	19620	95	154	
19506	49874	Α	19621	2	241	
19507	49875	Α	19622	63	320	RQWGNRLVTA/WSFFSWLSDC
		1				EGPGYGTIAEVGQLWGIMAKD
		l				LLRHSLASEELALNQNLTCHDQ
						FAQSTRSSARACGSIFCSTTQGS
19508	49876	A	19623	2309	2560	RNREARFYMTTPNFTETKR/WF
						HERVCALEREREREVTKISRLE
						RDNHRLVGDISQLKKELDQYL
		L_				QAISDLEDCNGKRLLQILEA

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19509	49877	A	19624	4994	7008	QWQKPKLTNDNLKFIWNQKR DRIAKTILSQKNKAGGITLPDFK LYYKATVTKTAWYWYQNRHI DQWNRTEPSETIPHIYNHLNFD KPDKNKKWGKDSLFNKWCWE NWLAICRKLKLDPFLTPYTKIN SRWIKHLNVRPKTIKILEENLGN TIQDIGMGKDFMTKTPKAIATK P\N*QMGSNILNGQKLEAFPLKT GTRQGCPLSPLLFNIVLEVLAR AIRQEKEIKGIQLGKEEVKLSLF ADDMIVYLENPIVSAQNLLKLIS NFSKVSGYKINVQKSQAFLYTN NRQTESQIMSELPFTIASKRIKY LGIQLTRDVKDLFKENYKPLLK EIKEDTNKWKNMPCSWVGRINI VKMAILPKVIYRFNAIPIKLPMA FFTELEKTTLKFIWNQKRARIA KSILSQKNKAGGITLPNFELYY KATVTKTAWYWYQNRYIDQW NRTEPSEITPHIYNYLIFDKPGK NKQWGKDSLFNKWCWENWLA ICRKLQLDPFLTPYTKINSRWIK DLNVRPKTIKTLEEDLGITIQDI GMGKDFMSKTPKAMATKDKID KWDLIKLKSFCTAKETTIRVNR QPTKWEKIFATYSFDKGLISRIY NELQQIYKKKTNNPIKKWAKD MNRHFSKEDIYAAKKHMKKYS SSLAIREMQIKTTMRYHLTPVR
						MATIKNSGNNRQ

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19510	49878	A	19625		764	GSSEKRKFKTGNQKLSMEVEK NQQMRSLSLYQQRVEGSKCAV RRKGGSPKPKAFLAAAEKFLN VPVMRSASHEQLVLTGDPNKT LPGSKSRLSGTDSPRACSTFPIW ATKIVTEMQVQLFAACRVQLIK HPRPSAARTAMEVLESGEQGV LQWDRKLSELSEPGDGEALMY HTACQGHEEQVWEEEALQLGL RSGILLNHWKRQLVGGAGILRS AQLHSPYHHWQMISLVSPSPAV MAVMENDDICMVLCVKVMDE PNHILDAISEHQIPSLEDQWFSE CALWTSKAYENLVERHILGPHC RPTELETLGISEDDIDHNPYLDF PGVAGLGSRRNVKERGWLSEM KKEGDIFMKIQG*AICPTSGPAE GLPPPTPVPHVPDRPCGT*GP/PP SPGSDSS\PACGPTAARPAPRSP APPWRCGPRWAEDVLLTGLCK QQIAEPAFRLQFWWPI

2457

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19511	49879	I <sub>A</sub>	19626	<u> </u>  1	1430	L MNGTAEGQHPWVHSTFNGEPQ
19311	49879	A	119020		1430	SLSSSVRVKNQGPRDQAELGFQ
		1			·	GNGSTVPAAQARTIARARGSGS
						GGGDALQGGGAGAGPGATKR
						TARLLARGLRRAALRPDGRGV
						LGAGAGSSQRRGGGRRKMETH
						ISCLFPELLAMIFGYLDVRDKG
						RAAQVCTAWRDAAYHKSVWR
						GVEAKLHLRRANPSLFPSLQAR
						GIRRVQILSLRRSLSYVIQGMAN
		ŀ				IESLNLSGCYNLTDNGLGHAFV
						QEIGSLRALNLSLCKQITDSSLG
						RIAQYLKGLEVLELGGCSNITN
						TGLLLIAWGLQRLKSLNLRSCR
						HLSDVGIGHLAGMTRSAAEGLP
1						GPGAAHATGLPEAHRSFSKAHL
						PRADGPEAPQPQLLWGNLGRW
						PPAPVAHGQPAHAQPCAPCDNI
						SDTGIMHLAMGSLRLSGLDVSF
		İ				CDKVGDQSLAYIAQGLDGLKS
						LSLCS/RRGLRRAALRPDGRGV
						LGAGAGSSQRRGGGRRKMETH
						ISCLFPELLAMIFGYLDVRDKG
						RAAQVCTAWRDAAYHKSVWR
						GVEAKLHLRRANPSLFPSLQAR
•		1				GIRRVQILSLRRSLSYVIQGMAN
						IESLNLSGCYNLTDNGLGHAF/R
						AGDRLPARSQPEPLQADH*QQP
		İ				GPHSPVPQGP/ARCWSWEVAAT
						SPTLAFCSSPGVCSASRALTSAA
						AATFRMW\DRAPGRHDAQRGG
						G\LPGPGAAHATGLPEAHRSFS
19512	49880	Α	19627	293	1520	TTMDGSHSAALKLQQLPPTSSS
						SAVSEASFSYKENLIGALLAIFG
						HLVVSIALNLQKYCHIRLAGSK
		l				DPRAYFKTKTWWLGLFLMLLG
						ELGVFASYAFAPLSLIVPLSAVS
						VIASAIIGIIFIKEKWKPKDFLRR
						YVLSFVGCGLAVRG\TYLLVTF
		l				APNSHEKMTGENVTRHLVSWP
						FLLYMLVEIILFCLLLYFYKEKN
	1					ANNIVVILLEVALLGSMTVVTV
						KAVAGMLVLSIQGNLQLDYPIF
						YVMFVCMVATAVYQAAFLSQ
						ASQMYDSSLIASVGYILSTTIAIT
	1					AGAIFYLDFIGEDVLHICMFAL
						GCLIAFLGVFLITRNRKKPIPFEP
						YISMDAMPGMQNMHDKGMTV
						QPELKASFSYGALENNDNISEIY
						APATLPVMQEEHGSRSASGVPY
L	<u> </u>	1	<u> </u>	<u>L</u>	<u> </u>	RVLEHTKKE

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19513	49881	A	19628	751	2034	VSSSPTSLGCMRQKEYRIPKKL TNMPFLGSQPSLLALRSCHPTG RTCSA/SQVALFCLAAHRVSQIG ALLAIFGHLVVSIALNLQVSFHI RLAGSKDPRAYFKTKTWWLGL FLMLLGELGVFASYAFAPLSLI VPLSAVSVIGKTIIGIIFIKEKWK PKDFLSKFVLSFVGCGLAVVGT YLLVTFAPNSHEKMTGENVTR HLVSWIILFCLLLYFYKEKNAN NIVVILLLVALLGKLTVVTVKA VAGMLVLSIQGNLQLDYPIFYV MFVCMVATAVYQAA*VSQAS QMYDSSLIASVGYILSTTIAITA GKGFYLDFIGEDVLHICMFALG *VLAFLGVFLITRNRKKPIPFEP YISMDAMPGKVNMHDKGMTV QPELKASFSYGALENNDNISEIY APATLPVMQEEHGSRSASGVPY
19514	49882	A	19629	19	933	QSARPSLKRARSQRGRPLPSRA LVFLFLFFFFFFNFLPIHTQPSAH KDMTTNAGPLHPYWPQHL\RL DNFVPN\DRPTWAYTGLGLFSV TGGLRSWTTWAVVQGRAPGL\ PIGDFGRATVPCCWVLQWCGFI HLVIEGLVPFSNYERPCFGDQSL LYSQLW\KEYGQGETARYILGV TNFTV\CM\ETITSFA\WGEPFSL WVVIV\FLLQHPL\RFIL\QLVVS VG\QIYG\DVLYFL*RATPTDFQ HGKPGPTPLLLPGFSFVFHEMP LWLVPAPGSPLCLDA/V*KHPQ LMPQSNAWNAKGPQKPKSKK
19515	49883	С	19630	113	257	
19516	49884	A	19631	143	330	HHGPVPFPRGLIS*HSLHCPCPT PGGKNVFFFCEYFLKLAGIETT NCSADKRGKKVCCLIPR
19517	49885	Α	19632	3	432	
19518	49886	С	19633	112	192	
19519	49887	A	19634		4410	MARPLALPLTGMSAPGSSLTTIP HTPSHPRSPSTPDTVSHPRSPSTP DTVSHPRSRYTPDTVPPTLPVY TRHRVPPTLPVYTRHRPTHAPG IHQTMCPTHAPGLHQTPCPTHA AGLHQTPSHPRSPSTPDTVSHPR SRSTPDTVPPTLPVYTRHRVPPT LTVYTRHRVPPTLPVYTRHCPT HAPGLHQTMCPTPRSRSLPDTV PPTLPVYTRHRVPPTLPVYTRD RVPPTLPVYTRHRVPTHAPRLHQ TPSHPRSR

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19520	49888	Α	19635	1	444	
19521	49889	A	19636	3	1017	SSRPVRPRPAARLSAMSSTQFN
	1					KGPSYGL\SAQ\VKNRLL\SKYD
						PQKE\AELRTW\IEGLTGLSIGPD
						FQKG\LKDG\TIL\CTLMNKLQP
						G\SVPKINRSMQN\WHQLENLS
						NFIK\AMVSYGMNP\VDLFEAN
						DLF\ESGNMT\QVQVSLLALGG KRPKTKGAAEGGLDIGVKYSE
		İ				KQERNFDDATMKAGQCVIG\L QMGT\NKCASQSGMTAYGTRR
						HLYDPKNHILPPMDHSTISLQM
						GT\NKCASQVG\MTA\PGTRRHI
						YEYQAGNPTSCDNFSM\SLQ\M
						GYTQGAQTQSGQVF\GPGRPDI
						*TPSTCPQGTI\ADGAPSGTG\D\
						CPDPGEVPEYPPYYQ\EEAGY
19522	49890	A	19637	1	479	MQGLRVSHSTFYTLIPEVTENP
	1,50,50					HRLDFHWFLSAQLERVASGTIL
						KKKESPPRPKVLGSARWDLPV/
						PPGWCNSTALQ*SPRSDQV*SE
						RGMEAFFLPMPSESVASVCCRF
						SPWTSQKPQLSTSRDRKLQGKR
						*WFSTGVWAS*DHS*ERRCSQC
						PKTTCW
19523	49891 -	Α	19638	2	501	FVPEERSSETALQRCSLPPGRRA
						APDPGLQGLRGSLGGSGSELES
						RKAKLRAAEGLRRKTTEINQRD
						TADRCTRGHH*SPGESPPKPSAF
						SL*LSVPRSGKLRFLRSPRREPA
						AN*SHRLRRHREKKCFHPPFTL
						NLIASGTSLEGGGVAPTWSTGR
						SQRALPSTL
19524	49892	Α	19639	$\lfloor 1 \rfloor$	1122	RFLILINHAVFEYQDIFVASFPSI
						FFNIHCRGEGEREKGFAGKDRT
						PPPSTLLCGSGHLPPLASRRPPR
						PRCACCLPGGPLSSSALEAG*Y
						WRFPKLSG*KRSHTPWVPQPQQ
						PLPNPSPGKWVWASSEALQ/GT
						RA*RGIAELTHLLRLPARRGTQ RPA\PGHRAGASPQFRREASSRG
						PGLRPRPGRTRELGFPKRRVWA
						TAASWPRGTEPCAAA*RRRRRR
						RRARGRELSPRSTWSSDTESTF
						ALSCDPS*PRHQSVPKGRVGSC
						KGKLANKGFFCPTWQRPIRVAS
						VDMELLHFVPKRALPGFTIAISF
						LSSAPPLCPWYSGSNPWLRNPL
						KPRLAALFQRSAAPRLHFSGAA
						,
	l			<u> </u>		SLQVGERRQIRDFRVSAGA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19525	49893	Α	19640	14	618	GNENVKSGRKTSLSPFVLSDFT
						ASSALLASDSSRWMSLQSASEC
						\PGRAAGSGPGYG*PVSEVTGG
		İ				PLSAHTHSSL*TSQRGCPPG\PE
						HHGAGPFP*GPGLKDVSGRGG
						NGCWGRRKPAHMGGGLGPAT
						PLQFPPRPSPPPPPLGPPQ\PPPPP
						SPPLGPASSPP/CLPHRPPPPPPPP
						PSRCVTAPCVTAPGPAPTGAAA
						RRH
19526	49894	В	19641	42	121	
19527	49895	A	19642	1	153	
19528	49896	A	19643	104	355	DDDTMAIL AICIAN LLTVI OVCD
19529	49897	A	19644	459	988	DPRTMNLAISIA\LLLTVLQVSR
						GQKVTS\LTA\CLVDQSLRLDCR HENTSS\SPSKYEFSLTRETK\KH
						\VLFGTVGVPEHT\YRSRTNF\TS
•				-		KYNMEGPLTYSAFH*ARTRGTL
						QVLQFHQFGALSPPIFLSRNV\T
					·	VVQRTNWVKC\EGI\SLLAQNTS
						WL\LRLLLS\LSLLQ\ATDFMSL
19530	49898	A	19645	1	950	WE LET CE EST CONTROL WITH THE EST
19531	49899	В	19646	55	1459	
19532	49900	A	19647	1	165	
19533	49901	Α	19648	183	700	PNGNQPPRETPQRISRGIQRLGT
						AVASSGGSRSLPRSASAEPAQG
						AAARRAAPGPAGRCRAARGAG
						AGSARRLP*APAHPAPPEQGRR/
						SRTARSRAAPQPPPPSQPQSPQL
		1				GPRPSQGAPQRRGPSSPGPRAG
						CRAQ*EALTPRLWRSSRTPSSR
		<u> </u>				GLGPHVRGGIFSPKIPT
19534	49902	В	19649	71	264	
19535	49903	Α	19650	3	810	GVSPCWPGWSRTPDFGSNPKCP
}						PIRASPGAELQALSSTVTTPYW
						GILVTAVFPH*GLRPRQCRQDH
		1				PAGRQGPGPGEVPEILGQSGCT
		1				DRTWSKAGGRTQAPGPRSRAG
						RRVSGQEIRAPGPLGCRHGG/V
						GAPWTPEAASPLTATEPSCPH/L
						QAPCGYMPLSVSPRRRYRGPA
						GDQKVKMLKFKAFCLDYWQF
						LCLQPLHGAYKRDSDLMTWIW
						GLLPEVTGAAGTTSPNVHTSGR
						FFRACVFCPVHTLVKKEPHPGQ
L	<u> </u>	<u></u>		L	l	QEIIMEPSPWSP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
1				sequence		
19536	49904	A	19651	258	1378	REILSLDRLPGATPNQALSAEA
1,7330	17704		17031	250	1370	ACPKGQTSLLCGAMGIQGLAK
						LIADVAPSAIRENDIKSY\FGRK
						VAIDASMSIYHFLIAVRQGGDV
						LQNEEGETTSHLMGMFYRTIR
		l				MK\ENGIKPI/YKSGELAKCSEL
	;					R\AEAEKQLQQAQAAGAEQVV
						EKFTKWLVKVTKQHNDKCKH
						LLSLMGIPYLDAPIEAEASCAAL
						VKAGKVYPVATEDMDCFTFGS
		l				PVLMQHLTASSDYCKSIWSIGP
						KRAVDLIQKHKSIEEIVQRLDPN
1						KYPLPENRLHKEAYQLFLEPEV
						LGPESVELKWSEPNEEELVKFV
						CGEKQFSEERIHSGVKRLRTEK
				•		GGGEPRGP/DWTQPGRASPTKR
						PDHQEDQHTRSRSSEGSH*GWL
19537	49905	A	19652	1040	2322	LPACPPASTPASTMSIMVTQKS
	•					YKVST/SRWSSGPQAFGNCSYA
		ŀ				RGPGACISSLSFSRVGSSIFRGVL
		ŀ				GGSYGGASGMGSITAVTVNQS
						LLSPLNLEVEPNIQAVCTQEK\T
			:			LNNKFASSIDKVQFLEQQNKML
						ET/ESYINSLKQQLETLGLERLK
						LEAELGNMPGLVEDFKNKYED
						EINKRTEMENEFVLIRQYVDEA
						YMNKVELESCLEGLTEEINLLG
						QLYEEEIQKLQSQILDTSVVLC
						MNNSHSLDMDSIIAEVKAQYEE
						MANHSQAEAESMYQIKYEELQ
						TLAGKHGDDLRHAKTEI/S/EM
						NQNGAKQDMA/*QLHEYQELT
						NIRLALDIKITIYRKLLEGEESQL
						ESGMQNMNIHMKTTSGYAGGL
						SSAYEGLTSPSLSYGLGSSFGSG
						MGSISFSHTSFTRAIVVKKIEIHD
						GKLVSESSDVLPK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19538	49906	A	19653	1040	2322	LPACPPASTPASTMSIMVTQKS
						YKVST/SRWSSGPQAFGNCSYA
						RGPGACISSLSFSRVGSSIFRGVL
						GGSYGGASGMGSITAVTVNQS
						LLSPLNLEVEPNIQAVCTQEKIT
						LNNKFASSIDKVQFLEQQNKML
						ET/ESYINSLKQQLETLGLERLK
						LEAELGNMPGLVEDFKNKYED
						EINKRTEMENEFVLIRQYVDEA
						YMNKVELESCLEGLTEEINLLG
						QLYEEEIQKLQSQILDTSVVLC
						MNNSHSLDMDSIIAEVKAQYEE
						MANHSQAEAESMYQIKYEELQ
						TLAGKHGDDLRHAKTEI/S/EM
						NQNGAKQDMA/*QLHEYQELT
						NIRLALDIKITIYRKLLEGEESQL
						ESGMQNMNIHMKTTSGYAGGL
į	,					SSAYEGLTSPSLSYGLGSSFGSG
					İ	MGSISFSHTSFTRAIVVKKIEIHD
						GKLVSESSDVLPK
19539	49907	Α	19654	152	207	
19540	49908	В	19655	75	402	
19541	49909	В	19656	1	2319	
19542	49910	В	19657	1	1341	
19543	49911	Α	19658	198	1118	LHAPNSLHHHLPRDLL*LLPLPL
						QHRVSDPGLHSALRPMEEHRA
						QS*QPSAPEDAVQA*WGHGGH
-						SPGPDRAGCHHCRGDQERVST
						HHVLPVCHHPDDAYGGCGAG
						WNPDL/PGQMRSHWMSPKIRPA
						NWTRTSWWALPQAPGLSPGAQ
						SWPSFVPRTTPTTPGTTCPTPSW
						*SWRSTSRTSSSLNPFTESLKNS
						LRTSKPFGWSQSAMATPCPLLP
						PASRVELWPETW\PLGQGHATS
						SQWKCVPERKL*QGGEAGGEQ
						LGREPKPSPPSSFLTGQRQEKSP
						EEYCSLLVPLQYFGSRGTHADL
						LHG

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19544	49912	A	19659	l	2707	MPGPHPASASCPPRLFTPRRRAF CPLRICAPLAAAAFCPPRFLPHT RRRAFLLLCRRGFLPPPCLGGG LAAAFFPHPPLPFLRPHRRGFLP HRRRGFLLPPRLGFLPAAAFYPP PPRVFARRGFVPPSPPRLFARRA FCHTPPAAAFSPSAAAAFCPTA DTAFCTLAALAFCRRGFLPPRC RGFLPLRHRRSFLPHRRRGFLPT AAFCHRGFLLATAFYPPRLFAP RCHGFLPPCRSDFFAAAAFCPR RRGFLPLSPPLFAPVTAAFCHRDFL PPPRIFVPAATALRAGAADSA ASSTGVLARAALRGPCGLHRV ALPVEAMGEKKEGGGGGDAA ATEGGTGAAASRALQQCGQLQ KLIVIFIGSLCGLCTKCAVSNDL TQQEIQTPEPAKPAKPASQTRQ PSQPSHPASQASQTSQPAKQPA SQASQPAKPAKPPSQPSQPSHR ASQASQATQPERPASQPASEPS QPSLPASQASQDRQPAKPAKPD SHPRQPSQP/YQPSQPASQASQA SQPAKPSRQPSQPRQPSQLASQ ASQSTQPLKPAKSPVQQSQPSQ PASPDSQASQASQATQPAKPAK PPSQAAQPAKPAKPPSQPSQSSPP
						PSQRAKPAKPARQPSQPSQSS ASQPSQACQPSQPGSQASQAS AAKPGMPAKPARQPSQPSQPA QARHASQAIQPAKPAG*PSQT

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19545	49913	A	19660	1223	2286	RQCFKATEQPPAIHHPSSQPSRQ PSSQPPIYPSPICGRKSGIQPKFQ LNG*MVNG*MDE\SRERHEWP CNYQVRPKAGEKQPGAEANRE TRPEGGTP/RGERGRQGPATRR QKSTPEAGPSDLEKQIARQEAP AGTDEIERQQARPH*AP*KIPVE WVNGEWLNGRMPGDPHEWQC NYQPAIQPAIQPANQLSIHNLW TQFRHSTKIPVEWVNGEWLNG RMPGKHHEWQCNYQPAIQPAI QPANQLSIHNLWTQFRHSTKIP VEWVNGEWLNGRMPEDHHEW QCNYQPAIQPAIQPASQLSIHN MWTQFRHSGQIPVEWVNGSRL HGRQQGIFMHGNATSSHSPSIQ PAIQPAIQPANQLSIHNLWTQFR HSTKIPVEWVNGEWLNGRMQR NIMNGNATISHSPSIQPAIQPAIQ PVNHLSIHNLWTQFRHSTKIPV EWVNGEWLNGRMPGKNHEW QCNYQPAIQPAIQPANQLSIHNL WTQFRHSTKIPVEWVNGEWLN GRMPGDHHEWQCNYQIAPML QSNRTAASHSPSIQPAIQAAIQS ATHLSIPNLWTQVRHSTKIPVE WVNGEWLNGRMPGRDMNGH ATIRYGPKPEKNNRAPRQTEKQ GQKGAHQRREGKARAGNEETE KYSRSRAIRPRKTDSASRSSRGH RRNRKAASAATLGPIEDTNQNK
19546	49914	A	19661	39	599	LHVRDTGLEVKAISIQGPGTLM GASPNQGQNRPHARQRAPPQ/G /PPGEPERRAPLPSGHGEPCRHR PPPFPQPP/AGTQKPLLQGPGGG *PAENAPTAALGSPAPPRGCQA APPPRSGAGRPDLPTLAGPRPAP A\PPPSAAPPPPSGAPSR/PAAG RQRLSGVSSGPSLGWW*VGRG RGLPAFAQIAGHQVGPRRRRTP AGRKPRSPAGPR
19547	49915	A	19662	39	599	GASPNQGQNRPHARQRAPPQ/G /PPGEPERRAPLPSGHGEPCRHR PPPFPQPP/AGTQKPLLQGPGGG *PAENAPTAALGSPAPPRGCQA APPPRSGAGRPDLPTLAGPRPAP A\PPPSAAPPPPPSGAPSR/PAAG RQRLSGVSSGPSLGWW*VGRG RGLPAFAQIAGHQVGPRRRRTP AGRKPRSPAGPR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19548	49916	A	19663	1	379	YFPHTAIRYPPHLNRQDPLKDL
						VSLACDPASQQPGPLNGSGQLK
						MPSHCLSAQMLAPPPPGLPRLA
						LPPATKAATTSEGGATSPTSP/S
						PGIWDSKGLLPSPLLHRPRNPR
						GQHSRPPAHVFGGKLE
19549	49917	Α	19664	3	285	
19550	49918	Α	19665	2	1556	RLEFPGRRFRSAPAPAGPAPPA
						APGMYSSPLCLTQDEFHPFIEAL
						LPHVRAFAYTWFNLQARKRKY
İ						FKKHEKRMSKDEERAVKDELL
						GEKPEVKQKWASRLLAKLRKD
		]		ļ		IRPECREDFVLSITGKKAPGCVL
						SNPDQKGKMRRIDCLRQADKV
						WRLDLVMVILFKGIPLESTDGE
						RLVKAAQCGHPVLCVQPHHIG
						VAVKELDLYLAYFVRERDAEQ
						SGSPRTGMGSDQEDSKPITLVT
	ļ					TDFQESFVTSGVFSVTELIQVSR
						TPVVTGTGPNFSLGELQGHLAY
						DLNPASTGLRRTLPSTSSSGSKR
		l		]		HKSGSMEEDVDTSPGGDYYTS
						PSSPTSSSRNWTEDMEGGISSPV
						KKTEMDKSPFNSPSPQDSPRLSS
1						FTQHHRPVIAVHSGIARSPHPSS
						ALHFPTTSILPQTASTYFPHTAIR
		İ				YPPHLNPQDPLKDLVSLACDPA
						SQQPGPSWYLG*QRSSSLAPSPS
						SQESQGAAQPAPGPRFRWKIRV
						NKNTPADSQPGQKDKTHRRTH
						SGGKEKTKAE
19551	49919	Α	19666	1	477	
19552	49920	Α	19667	1	440	WTQDM/EGGISS/PVKKTEMAK
						SPFNSPSPQDSPRLSSFTQHHRP
						VIAVHSGIARSPHPSSALHFPTT
						SILP\QTASTYFPHTAIR\YPPHL
						NPQDPLK\DLVS\LACDP\SSQQP
						GP/SYSPPDTSPANR\SFVGLGPR
				İ		DPAGIYQAQSWYLG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence	ŀ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19553	49921	Α	19668	2	1963	CPTGFEHGSPCARRTPTGRVSV
						TRCRRRAWGSLRVSVCGPRAQ
						LQAELVQEEQGGGVLCVIVMP
						LTPGRTPRLPRGRALCNGRGGV
						WGGGGSPRGLLPAQAPGDPEV
						CKFRFLDRCFPPYAGPSDLSLR
						GRAVGPPSASLALLTSRLSGLQ
						DEFHPFIEALLPHVRAFAYTWF
						NLQARKRKYFKKHEKRMSKDE
						ERAVKDELLGEKPEVKQKWAS
						RLLAKLRKDIRPECREDFVLSIT
		ŀ				GKKAPGCVLSNPDQKGKMRRI
						D\CFRQADKVWRLDLVMVILF
						KGIPLESTDGERLVKAAQCGHP
						VL/CALKPQHI\GVAVK/ENLDL
				:		YLAYFVRERDA*QSGSPRTGM
						GSDQEDSKPITLDTTDFQESFVT
						SGVFSVTELIQVSRTPVVTGTGP
						NFSLGELHAHLAYDLKPASAGL
						TRTLSSTFSSGNKRY\KAGSMEE
						YVDSTPWGDD*T*PSSPTSSSRN
						WTEDMEGGISSPVKKTEMDKS
						PFNSPSPQDSPRLSSFTQHHRPVI
						AVHSGIARSPHPSSALHFPT\TSI
						LTPDGPSPTFPHTAIRYPPHLNP
						QDPLKDLVSLACDPA\SQQP\GP
						LNGSGQLKMPSHCLSAQMLAP
						PPPG\LPRLALPPATKPATTSEG
						GATSPTSPSYSPPDTSPANRSFV
						GLGPRDPAGIYQAQSWYLG
19554	49922	A	19669	332	551	ESWLGRGHQPPP\PGLFHSPKPS
.,,,,,						SGTAVLPAWRPPSPGPGAGDD
						ASPSPS*AGPGAGANPGA*QRS
						ASGEKGS
19555	49923	Α	19670	1614	1673	
19556	49924	Α	19671	72	227	EEFPMGHTA*WSCEKRATILQI
						QEW*IQQQLAPCAWQSHRHST
						RAHESSHG
19557	49925	С	19672	1	604	
19558	49926	Α	19673	653	889	VYPQTNWKQNTGTRVFIAALF
						TM\AKGENK*PSMGEW/IKQM
						WSIHTMERHAAIKRIEAQVYAT
						M*MKLKSMLLLERNQT

SEQ ID	-		SEQ ID NO:	L		Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/340,217	sequence	or peptide sequence	deterion, (-possible nucleotide insertion)
					•	
19559	49927	Α	19674	75	1619	EVSIPAQTMWTADEIAQLCYEH
						YG\IRLPKKGKPEPSHEWTLLA
						AVVKIQSPA\DKACDTPDKPVQ
						VTKEVVSMGTGTKCIGQSK/M*
						RKNGDIL\NDSHA\EVIA\RRSFQ
						RYLLHQLQLAATLKEDSIFVPG
						TQKGVWKLRRD\LINVFFSSHT
						PCGDASIIPMLEFEDQPCCPVFR
						NWANNSSVEASSNLEAPGNER
1						KCEDPDSPVTKKMRLEPGTAA
						REVTNGAAHHQSFGKQKSGPIS
						PGIHSCDLTVEGLATVTRIAPGS
						AKVTDVYRTGAKCVPGEAGDS
						GKPGAAFHQVG\LLRVKPGRG
						DRTRSMSCSDKMAR\WTVLGC
						QGALLMHLLEEPIYLSAVVIGK
						CPYSQEAMQ\RA\LIGRCQKV\S
	İ					ALPKGFGVQELKILQSDLLFEQ
						SRSAVQAKRADSPGRLVPCG\A
1						AISWSAVPEQPLDVTANGFPQG
					,	TTKKTIGSLQARSQISKVELFRS
İ						FQKLLSRIARDKWPHSLRVQKL
						DTYQEYKEAASSYQEAWSTLR
L		L				KQVFGSWIRNPPDYHQFK
19560	49928	C	19675	45	140	+ WW. TO WOOD WOOD AD TO SO
19561	49929	Α	19676	3	354	AKIKTGYK\REHINLGCDMDFDI
						VGPSIRGALVLGYEGWLAGYQ
						MNFETAKSRVTQSNFAVGYKT
						DEFQLHTNVLLLNLERTCFLTG
						PTPTPPPLSWLPGTTFSLTWAM TGRYTINSQ
19562	49930	A	19677	 	983	AARSAPLPGSAPCLRVAAPAVA
19302	49930	Α.	19077	1 1	763	ASEPAATSSEQKMAVPPTYADL
				ļ		GKSARDVFTKGYGFGLIKLDLK
	]					TKSENGLEFTSSGSANTETTKV
						TGSLETKYRWTEYG\LTFTEKW
						NTDNTLGT\EITV\EDQL\ARGT*
1						SLTF\DSSFSP\NT\GKKNAKIKT
						GYKREHINLGCDMDFDIAGPSI
						RGALVLGYEGWLAGYQMNF*
						DCKIPE*PQSNFWQLANKT\DEF
						QASR*M*ND\GTEFGGSIYQKV
						NKK\LETAVNLPWT\AGNSNTR
						FEIAAKYQ\VDP\DACFSGKVNN
						SSLIGLGYT\QTLKPGIKLTLSAL
						\LDGKN\VNA\GGHKL\GLGLEF
						QA
L	<u> </u>	<u> </u>	<u> </u>	L	l	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19563	49931	A	19678	551	831	CHPSSHVLCAPASLSDHQWPPP ALCPSSRPQ/PCALPWSGTLSSR PQPCPTHPSGIEA*LCLPLVTQL G*LPL\PAPQHLVPPCPSTCPTM LKLF
19564	49932	A	19679	47	980	VCQYCTARMADFGISAGQFVA VVWDKSSPVEALKGLVDKLQA LTGNEGRVSVENIKQLLQCLVP GSTT\LHSA\EILA\EIARILRPGG CLFLKEPVETAVDNNSKVKTAS KL\CSALTLS\GLVEEKELQREP LTPEEVQSVREHLGHESDNLLF VQITGKKPNFEVGSSMQLKLSI TKKSSPSVKPAVDPAAAKLWT LSANDMEDDSMDLIDSDELLDP EDLKKPDPAS\LRA\ASCGEGKK RKACKNCTCGLAEEL\EKEKSR EQMSSQPKSA\CGNCYRGAMPS GCASCPYLGMPAFKPGEKVLLS DSNLHDA
19565	49933	A	19680	29	1270	FPFWPAVFQVCQYCTARMADF GISAGQFVAVVWDKSSPV\EAL KGL\VDKLQAF/TPGNEGRVSV\ ENIKAAVAILPTKNPSFGHYFV Q/CLVPGKAPLWHSA*DFWAG NPPGFLRPGWMFFFLKEPVETA VR*Q*AKWKTASKL\CSAL\TLS GLV/EKLKELQREPLTPEEVQSV REHLGHESDNLLFVQITGKKPN FEVGSSRQLKLSITKKSSPSVKP AVDPAAAKL\WTLSANDMEDD SMCIFCGCSLTHRWPLEHVVQV E/IMMDQPKRRTRVDT\FFTPRT PKFPSRSPASHFSFSIKQKT/TRP VSLIALNTL\QDLIDSDELLDPE DLKKPDPSLLRAASCGEGKKR KACKNCTCGLAEELEKEKSRE QMSSQPKSACGNCYLGDAFRC ASCPYLGMPAFKPG\EKVLLSD
19566	49934	A	19681	3	405	QDFGTRALAAVQIKNGGSWWC PPVSLAAGVGPRPQLGRPYVRV APHGQV\GLSFAMWKTLT\FFR GASPGVGSQAMLKCVA*SPHH WRARRDPEFIAYP\HLR\IRTKP\ FPWG\DGNHTQQYITPH\VNPLS NLATED

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	•	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19567	49935	A	19682	2	450	VNKAGGLIYQLDSYAP/RAEAE
						KTFSYPLDLLLKLHDERVLVAF
						GQRDGIRVGHAVLAINGMDVN
						GRYTADGKEVLEYLGNPANYP
						VSIRFGRPRLTSNEKLMLASMF
						HS\IKFVVLADP\RQAGIDSLLRK
						IYEIYSDFALKNPFYSLEMP
19568	49936	Α	19683	1	2034	
19569	49937	Α	19684	202	453	RRLWRGSCPRCGGKARDGMG
						VRGR/YSEG/RERV*FGGPPRGS
						TWPLCGRWRRPARPSAPLRWS
			<u> </u>			AMCPNTSDAAGACAAAVRCG
19570	49938	Α	19685	269	326	
19571	49939	В	19686	98	562	
19572	49940	Α	19687	1	2223	
19573	49941	Α	19688	374	1107	IPVSERQGSDWRFFSVYVVNKA
						GGLIYQLDSYAPTGLRLEKTF\S
						YPLDLAGSS*HDERVLVAFGQR
						GRHPEWGHAVL\AINGM\DVNG
			}			RYTADG\KEVLEYLG*PLLNYP
						VSIRF\GRPRL\TSNE\KLMLGLP
			,			CFHSLFAIG\SQACLPGTREASR
			ļ			HLKILGRQDTFK\LH\CYQTLT\G
-						IKFVVL\ADPR\QAGI\DSLLRKI
						YE\IY\SDFALKNP\F\YSLEMPIR
			İ			CELFDQNLKLALEVAEKAGTF
19574	49942	Α	19689	119	288	ALHLNSSDVHQKLLPVQSLKKS
						*GRHRNNTCNQRVHA/SSTSRE
						FFTKLNKDNDILKP
19575	49943	Α	19690	92	317	KSVGPVNFPLPSGNMNEDN*M
						H*HPFSSVTQAFRLRVLQSII\QR
						PL*TFNPGSSSLFSSISGQSNFPL
						GVAVLRP
19576	49944	Α	19691	27	389	NFKTLKNN*IYK*MCSQEIKILS
						*KLIRFWYEIHAFFPWAHGVVF
		1				CVAQEFCLPTVVLLPVPRLLIM
		1				KDMVRRLQELRHTEQVQRAYA
						LNCGEGATVSYEIQIRVLREFGL
						ADAAAELLQN
			I			ADAAAELLQN

19577 49945 A 19692 313 2255 KTPQVQPFPRQTGAAGAGAP SGRSGERAAGRAGPSGGSGGG QDSAHAGEAYRVRGILLRKTFQ CLKDYFLTYYTSSKVVPVKVPC RQVLTIDKCSLPRSSHRNARTV SNDFPRKSCSRERCTHCMRPPE GRRPPNGMNSWKWPSGDESK WRRGEPGAVPEPEPLTGEERAA AAAVGGGWPGVLAPVRTAVL VAAAAEATAAVPAAAAAAGG ASGHCNHPVMQGRNLSHRFLL PLHHSPYQFGTSSYSQQYGCE SKLYSLDHGHEKPQDKKKRTS GLATLKKKFIKRRKSNRSADHA KQMRELLSGWDVRDVNALVE EYEGTSALKELSLQASLARPEA RTLQKDMADLYEYKYCTDVDL IFQETCFPVHRAILAARCPFFK/I TAFFLTRVWGRDNNGHQYSW Y*YAHVFCFVTLPLYRRVWEW EDSRFQNVDILVQLSEGIWEHQI PLM*DMRGTLLI/DMCYYDVVL SFSSDSELVEAFGGNQNCLDEE LKAHKAVISARSPFFRNLQRRI RTGEEITDRTLRTPTRIILDESIIP KKYATVILHCMYTDVVDLSVL HCSPSVGSLSEVQALVAGKPN MTRAEEAMELYHIALFLEFNM LAQGSCWECIPPNDEANQEQRS EGSRKWAPTREGSKGRSHASA	SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
EGSRKWAPTREGSKGRSHASA	19577	49945	A	19692	313	2255	SGRSGERAAGRAGPSGSGGG QDSAHAGEAYRVRGILLRKTFQ CLKDYFLTYYTSSKVVPVKVPC RQVLTIDKCSLPRSSHRNARTV SNDFPRKSCSRERCTHCMRPPE GRRPPNGNMNSWKWPSGDESK WRRGEPGAVPEPEPLTGEERAA AAAVGGGWPGVLAPVRTAVL VAAAAEATAAVPAAAAAAGG ASGHCNHPVMQGRNLSHRFLL PLHHSPYQFGTSSYSQQGYGCE SKLYSLDHGHEKPQDKKKRTS GLATLKKKFIKRRKSNRSADHA KQMRELLSGWDVRDVNALVE EYEGTSALKELSLQASLARPEA RTLQKDMADLYEYKYCTDVDL IFQETCFPVHRAILAARCPFFK/I TAFFLTRVWGRDNNGHQYSW Y*YAHVFCFVTLPLYRRVWEW EDSRFQNVDILVQLSEGIWEHQI PLM*DMRGTLLI/DMCYYDVVL SFSSDSELVEAFGGNQNCLDEE LKAHKAVISARSPFFRNLLQRRI RTGEEITDRTLRTPTRIILDESIIP KKYATVILHCMYTDVVDLSVL HCSPSVGSLSEVQALVAGKPN MTRAEEAMELYHIALFLEFNM
11057X 1/100/16	19578	49946	В	19693	4	3923	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19579	49947	A	19694	1	2154	MEYYAAMKTKEIMSFALTWIE
1,3,7	',	ļ ` `	1203.	1		LEAIVLCELTHVLSELMQKQKT
						KYHMLSLGSPVPTFALWNFDV
						AQDFHHFIENARSSQCRGRPRM
						TGAGAGAPSGRSGERAAGRAG
						PSGGSGGGQDSAHAGEAYRVR
			:			•
		ŀ				GILLRKTFQCLKDYFLTYYTSS
İ						KVVPVKVPCRQVLTIDKCSLPR
		İ				SSHRNARTVSNDFPRKSCSRER
ŀ						CTHCMRPPEGRRPPNGNMNSW
İ						KWPSGDESKWRRGEPGAVPEP
ŀ						EPLTGEERAAAAAVGGGWPGV
-						LAPVRTAVLVAAAAEATAAVP
						AAAAAAGGASGHCNHPVMQG
						RNLSHRFLLPLHHSPYQFGTSS
						YSQQGYGCESKLYSLDHGHEK
•						PQDKKKRTSGLATLKKKFIKRR
·						KSNRSADHAKQMRELLSGWD
İ						VRDVNALVEEYEGTSALKELSL
						QASLARPEARTLQKDMADLYE
						YKYCTDVDLIFQETCFPVHRAI
ŀ						LAARCPFFK/ITAFFLTRVWGRD
İ						NNGHQYSWY*YAHVFCFVTLP
						LYRRVW\KEDSRFQNVDILVQL
						SEEFGTPNSLDVDMRGLFDYM
						CYYDVVLSFSSDSELVEAFGGN
						QNCLDEELKAHKAVISARSPFF
						RNLLQRRIRTGEEITDRTLRTPT
						RIILDESIIPKKYATVILHCMYTD
		ŀ				VVDLSVLHCSPSVGSLSEVQAL
l						VAGKPNMTRAEEAMELYHIAL
					,	FLEFNMLAQGSCWECIPPNDEA
19580	49948	Α	19695	1	163	
19581	49949	Α	19696	1	207	
19582	49950	Α	19697	1	579	PTRPGQAGSSSAMAAQRLAKR
						VL\SKLQSPSRARG/PCAGSPGG
						LQK/RGHARVTVKYDRPGSLQR
						RLDVEKWIDGPLEGLYRR\QEA
						D\MPMRSNIDELLE\LESEE\ERS
						RKIQGV/LRDS*KSCGKPVEDFI\
						QELLAKLQGLHRQ\PGLRQPSPS
						HDGSLSPLQ\DRARTASPLTLLA
						LFPGPPERRPALLCVLSCI
19583	49951	A	19698	2	151	PPSPPPLPPPPLPPAPSLPLSLPSF
						DLPQPPVLDTLGV/LF*LCRKKI
						SVS
19584	49952	Α	19699	2	310	
		لستت		<u> </u>		

NO:	of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19585	49953	A	19700		688	LTWMTERVDFTALLPLEPPLPP GTLPQPSPTPPDLEAMASLLKK ELEQMEDFFLDAPLLPPPSPPPL PPPPLPPAPSLPLSLPSFDLPQPP VLDTLDLLAIYCRNEAGQE\EV G\MPPLPPPQQAPSFLLPP\QPSR WGPLTHIPGQPTRGGPAKQKEE RPRNKSAALKVTPSGKRARG*S PWRAKSQGLEGTEFASLKERA\ ESVEREIQ\YVKD\LLIEVYKARS
19586	49954	A	19701	3	902	QRTRSC  AAMATPARPGEAEDAAERPLQ DEPAAAAAGPGKGRFLVRICFQ GDEGACPTRDFVVGALILRSIG MDPSDIYAVIQIPGSREFDVSFR SAEKLALFLRVYEEKREQEDC WETFGGAGGEASPA*RRSSSS GTRRWTWRTL*LGSSATATCW PC\GESDRQVLDLD/LGSTNARS SCARGRAGSGTCQGPSSWGPR G/CYSWYKGQPKTCFKCGSRTH MSGSCTQDRCFRCGEEGHLSPY CRKGIVCNLCGKRGHAFAQCP KAVHNSVAAQLTGVAGHLNTR LPARVNTQASLSLLKCPKLFF
19587 19588	49956	A	19702 19703	1359	1981	QSPFNDTLPGVPVSA*LHSLPPE HHPSKG*GVAKSPPVYPAGILL VCNNCAAYRKLLEAQTPSVRK WALRRQNEPLEVRLQRLERER TA\KKSRRDNETPEEREETDEQ RARRLQRDREAMRLKRANETP EKRQARLIREREAKRLKRRLEK MD\MMLRAQFGQDPSAMAAL AAEMNFFQLPV\NGVEL\DSQLL GKMAFE\EQNSSSLH

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19589	49957	A	19704	1	673	MSAKDGTCTLQDCMYKDVQK
						DWPGYSEGDQQLLKRVLVRAH
		1				NTQRQRQNERSDTRPCQRAAP
						RRFSHGTRRGTLPLASPDRARA
	1	ł				PRTGHCRTQRRLDRTGGGFPPL
		}				TGAWAMLADWIHEGTPSRGRS
						DNDEPSAGSVGGLRQHGPGPR
						APKLTTTKVKALASPVLGQKIA
						ALCLGLTAQGLNIMDNPGSLAI
						HVHPDSVLTRKTNPCTHIWGFT
						QREPPLVANMSAKDGTCTLQD
						CMYKDVQKDWPGYSEGDQQL
						LKRVLVRAHNTQRQRQNERSD
						TRPCQRAAPRRFSHGTRRGTLP
						LASPDRARAPRTGHCRTQRRLD
		ļ				RTGGGFPPLTGAWAMLADWIH
						EGTPSRGRSDNDEPSAGSVGGL
		1				R/HAWPRTKSPKANNH*SESTG
						QPGAGPKNCSPVLGAYSPGPQH
						HGQPGLPGHTCP/SRTQC*PGRP
						TPAPTSGVSLRGNPLWWPT
19590	49958	Α	19705	1088	2273	SPCLLTGGHLPADTSYRRAPAG
		İ		:		IWQVPLWDEASQRKE*YQHQQ
				İ		KKMSTQKPHPKDHNSSPAREQ/
						SSMENKFHELTEVGFRKITSLE
						KNINDLMELKNTARELREAYTS
						IKSRIDQAKESISEIEDQLNEIKC
						EDKVREKRMKSNKQNFQEIWD
						YVKRPNLRLTGVPESDRENGTK
			1			LENTLQEIIQENFPNLARQANIQ
						IQEIQRTAQRQSSSRATPIHIIVR
						FAKFEIKEKMLRAAREKDISMR
						QKINKDIQDFNTVLDQVDLIHV
						YRTVHPKSSEYTFFSAPHHTYC
						KTGHIIGSKTLLSKCKRMEIITN
						TLSDYSAIKLELRIKKLTQNHTT
					_	TWKVNTLLVNEY

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	l	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19591	49959	Α	19706	318	873	AYELTPEPSPLPVFPVSADGSTT
						HPLADKRKASPVPEQHEEHLPA
						RPVTQRCLTYPFPNITTASEFGP
						ATGFRARRSGRGSTVCCNPGCY
						PRARPEPEPRGTPPSVSAPRAAQ
						EAPARQR\AGPVLADTVP/SPAP
						RGKRDISPLFGNAHSASPKPSLD
						LWAPPP*AAGRRGPQQA*DVTE
						DTQQQWSAMAVGGTAKWM
						AALGRFLVTYVEVEGVKLQTF
	:					TVTTLTQSFKYSLSKYLQQQPL
						LSSRTIYPTFYYLSLLRNLRAFK
						LNAAEAKPMSSPQNLVLFLCFL
						SLQMAAPPIRLQTKGKQAQCQS
						STKSTFQRGRSLRGVSPTLSPTS
						QQPRNLDQQLASEQGEVGEAP
		1				LCAVIQGVIPVLGRSRSREEHPP
		1				VSALPGLHRKRQRDKEAGPVL
						ADTVPFRPHAGKGTFLRCLETL
						TRRAPNPRWTFGLPLRKLQGA
						VAHSRPEIRTHSVPNRT
19592	49960	Α	19707	1	690	DQDMTSSDIPVRLEGSLWLP*IF
						IGTFKAFDKHMNLILCDCDEFR
						KIK*GVDCGGRRLEDGNGLSG
						WAKQ*SLVSMTVEGPPPKDVS
		İ				QSRKP/CWRRGGPTVSQALC*IL
						YPSWGSHAPGSCRTCWASPWG
						WRAIPTGEELGKGYHLLVAKIK
						DKGEGP*FSNPVPTWPWGSSPT
1						YGPRSTPSR*GAHKETSAN**K
				}		MP\SWINDEI*PPMGMPPPGMRP
						PPPGMRGKCLQW

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:		Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ĺ	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19593	49961	Α	19708	1	1348	MVFWARPRTLLLCAALGLGAL
						YTATPAPTVAKRGQGTAQALS
						SEGASPKPWWLRHGIEPCKQSY
						WEVRTEQSPGVSERKAAAPVR
						GLEIKLPSPWDKAPEGGGGCGH
ĺ						SFSRLKCSFMPVLKRAVDLPAQ
						RSSPANGQTACSSGSLAPVLPD
						WEIPPSKGRQTPYATGELRLAS
						DGCPSGTKLPEEGTDSNLCCSA
						SSAESNNININININKKDAHGKT
						PSKGPQHQRSKLDKSMKTRKN
						HHKNAENSKNLNASSPPKDHN
						SSTAKEQNWTENEFDELTEVGF
						RRRVIANSSKLKEHVLTQCKEA
		•	<u> </u>		İ	ENFEKILEELLTRITSLEKNINDL
			<u> </u>			MELKNTAQEHREAYTSINSQID
						QAEERISEIEDKLNEIKREDKIR
		İ				GKRIKRNKK\SLQEIWDYVKRA
						NL*LIAVVPLNKITSPRTSETLST
		1				STNMSRSTLSCLLSSFSCPASCT
						ATSAALF
19594	49962	Α	19709	718	1138	GGRAAEGSQPPVAEPRCSRGAR
						ALPLPDTSWRQPS/APQAAP*D
						VELPASPSAPFAVLLSKPRLSQN
}						KPKSPFSVASP*HGDKRPQPSRH
						LI*IPHLSSRAPVHS/PLGWPSNR
			1	1		K/HKHLRPDTAHAPAEGPPLNY
10.50.5	100.60			<u> </u>	7.00	VRKQQKQS
19595	49963	Α	19710	5	760	RAAACGGAAPGRKRCAPGRKR
						CGPRRKRSLGPPTLGQRPRQPS
			1			QQLRKPRRRRGSPGSLPKSSPLP
						SSTGKTGWSTRGRSVRPPVTTS
						RCCPPGHPCSRRAPSCTARAPLP PT*PPTTAPCRASPPAPRPPRRPR
			]			PPPGTRSPTSRGRCRWTRSGAG
						TSRCSWRAGPGTAPAAAGARS
		ŀ				RPGAPGVSAAVPCAAACSWMT
				:		SAPEVRAASSPGTPSRGSGGRS
						HPPCSRGLPTTGPATTAPEAPG
						CWSCRRS
19596	49964	В	19711	180	300	o John de
19597	49965	A	19712	249	700	RQDPVSGGQSCPASCPSLWAA
			-			ARVWLGQ\PAPRAR*CPVCSSC
						PGPGAPEDTPEAGGCGCSGCRC
						CLRTQEPPAPLQPPHPPMAQTP
						REEVGRRGWTPGCPGTDPCLW
			[			GLRLLCPLWDPHRSGTRPWAQ
						VQRDTTETKAQRGLVRCQRGC
19598	49966	С	19713	273	485	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19599	49967	Α	19714	35	779	VRPPSHVTADSGRSPLSLTYLPL
		1				QEPGDMAAAVPRAAFLSPLLPL
		1				LLGFLLLSAPHGGSGLHTKGAL
		1				PLDTVTFY/KDYGDKLNMELSE
						KYKLDKESYPVFYLFRDGDFEN
		1				PVPYTGAVKVGAIQRWLKGQG
		1				V\YLGMPGCLPVYDALAGEFIR
						ASGVEARQALLKQGQDNLSSV
		1				KETQKKWAEQYLKIMGKILDQ
		1				GEDFPASEMTRIARLIEKNKRSC
		1				KKAVCDFPGFGGGREGRVNLL
		ļ <u>.</u>				AVSPLWNIRG
19600	49968	Α	19715	2	253	
19601	49969	Α	19716	494	1443	LAVPRPAYGPRSPGSLLRIPVPG
		İ		; ;		LHLLNLNVWSRGSAAALHTKG
						ALPLDTVTFYKVIPKSKFVLVK
	-	İ				FDTQYPYGEKQDEFKRFAENSA
		ŧ	ŀ			SS\DDLL\VAEVGISDYGDKLNM
		ŀ	<u> </u>			\ELS\EKYKL\DKESYPVF\YLFR
						DGDF\EEPSSHYTGGS*RLGAIQ
						RWŁKGGKGVYLGDGLELVCT V*LTPLAGEF\IR\ASGVEARQAP
						LEAGAQDNLSKCERRLRKKW
	1					AEQYLKIMGKILD\QGEDFP\AS
						EMTTESLGLIEKN\KMSDGKKE
	1					EL\QKSLN\ILTALPRRKGAEKR
						GAVNKAVCDFPRVWWEYGRG
						ELTLLAVSPLWNIRG
19602	49970	A	19717	286	332	ННННРНҮННННРНҮННН
17002	1.33.0	1				HPQYHH\HHPHYDHHRHRQGD
						HHGQLQYDDDHQPQCHQHDH
						LRCEHQHYIQ*WLINNLRASRY
						KPLLNIMLVFAAEMIVLVALGL
						MVVVVLELTVVVALTMTVVVI
						VRMMGGDIGDDGGGGSGDDG
						GGGSGDDGGGVMWQFFDSMS
			-			KNDSIIPTTTTTIIPTTTTTTIRA
						EQFTILTWKPVVSSQKNTSSFR
19603	49971	С	19718	.1	546	
19604	49972	Α	19719	1	90	
19605	49973	В	19720	217	951	
19606	49974	В	19721	67	1708	
19607	49975	В	19722	218	3267	

SEQ ID NO:	of peptide	Met hod	SEQ ID NO:	location of first	codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19608	49976	A	19723	1	634	LGRLSFPRPEQPDPGHRAHPSSP PRPRRPHARGRRGPRRHH/YHD
		1				TGANRAPEVRPPCPES/SPVGVA
						AFEPNGLRGQQPSPRSPGAPKG
						L*PTPPSASSSPKPREPSALGRPC
						CRCEESRFLEHRNSNPRVKDSR
						QARGRHAEFHGRTGVFERRRS
						RDHLPTHFPASVRWEKLSWCA
						RKNGLSGLRLCHGSQHFSGYSN
						RKVGTESLNQVWE
19609	49977	A	19724	3	1090	
19610 19611	49978 49979	A	19725 19726	23	636	FSYLPPGPSHGTWGLWELQFK
19011	149979	A	19/20	23	030	MRFGVCRHLMEDSMDMDVSP
						LR/PQNYLFSCELKADKDDHFK
						VDNDENEHQLSLRTPTVSLGVF
						EITPPVLLWLKCGSGPVHISGQ
						HLVAVEEDAESEDEEEEDVKLL
		1				RISGKTKTFMATNGKEYKHYKI
		1				SSEKSLDNKYKTRTPGFQAFGF
		1		-		EDLHPWPLGSQAFYLSLRVTPP
		1				VFLVLRLLDFD
19612	49980	Α	19727	1	830	MEDSMDMDMSPLRPQNYLFGC
						ELKADKDYHFKVDNDENEHQL
						SLRTVSLGAGAKDELHIVEAEA
						MNYEGSPIKVTLATLKMSVQPT
						VSLGGFEITPPVVLRLKCGSGPV
						HISGQHLVAVEEDAESEDEEEE
ŀ						DVNRVLFMKP/KRQGPQAYVR
						GFGVQLEIGSH*DPEQQF*RDSA
						QPSLATPSTPHLQALLCLSGQV
						LRLPLWGQRL*GLQGLLPPQHP
						EEHGVHVSPGQELHHQQGDPE
						PLPVLPTAEVL*SGHVQGVCEK
19613	49981	A	19728	1	684	RPKQEEEGGAQARVL
19614	49982	A	19729	3	535	DSVLRGCSLEQRSFISVRLLSYL
						SACRHPMEDSMDMDMSPLRPQ
			Ì			NYLFGCELKADKDYHFKVDND
						ENEHQLSLRTVSLGAGAKDEL
		1				HIVEAEAMNYEGSPIKVTLATL
						KMSVQPTVSLGGFEITPPVVLR
						LKCGSGPVHISGQHLVVYRRK
						HQELQAMQM\DCRALSTS*ASS
10615	140002	I.	10720	1	2260	APRPS
19615 19616	49983 49984	B A	19730 19731	1198	2268 472	AHDIGCPEVSPLVLEVARDEGP
19010	147704	\\\^	19/31	130	7/2	AHRPRFSH*SSHCSHWGTSQRA
						APPPCHPCIQTTW/PGAVTGLG
						G*PKTRSGPGKDRSSGAASSFP
	1					GGRV
L		Щ	L		I	OGICY

SEQ ID NO:	SEQ ID NO: of peptide sequence	ł .	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19617	49985	A	19732	92	2049	LTALPPAVSPGGHSSSSSSRTIS GIT*ELQR/LLQAVVPGPWQED VADAEECAGRCGPLMDCRAFH YNVSSHGCQLLPWTQHSPHTR LRHSGRCDLFQEKGEWGYMPT LRNGLEENFCRNPDGDPGGPW CHTTDPAVRFQSCGIKSCRVAA CVWCNGEEYRGAVDRTESGRE CQRWDLQHPHQHPFEPGKFLD QGLDDNYCRNPDGSERPWCYT TDPQIEREFCDLPRCGSEAQPRQ EATSVSCFRGKGEGYRGTANTT TAGVPCQRWDAQIPHQHRFTPE KYACKDLRENFCRNPDGSEAP WCFTLRPGMRVGFCYQIRRCT DDVRPQDCYHGAGEQYRGTVS KTRKGVQCQR/WSLE/TG*DYP ATTSLPLPPRFTFTSEPHAQLEE NFC/RDPDGDSHGPWCYTMDP RTPFDYCALRRCDQVQFEKCG KRVDRLDQRCSKLRVAGGHPG NSPWTVSLRNWQGQHFCGGSL VKEQWILTARQCFSSCHMPLTG YEVWLGTLFQNPQHGEPGLQR VPVAKMLCGPSGSQLVLLKLE RYVDNLGGWTKCEIAGRGETK
						GTGNDTVLNVALLNVISNQEC NIKHRGHVRESEMCTEGLLAPV GACEGDYGGPLACFTHNCWVL KGIRIPNRVCARSRWPAVFTRV

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19618	49986	A	19733		2296	HSAGQRSPSNDFQVLRGTELQH LLHAVVPGPWQEDVADAEECA GRCGPLLLSRAFHYNVSSHGCQ LLPWTQHSPHSRLWHSGRCDLF QEKGEWGYMPTLRNGLEENFC RNPDGDPGGPWCHTTDPAVRF QSCGIKSCRVAACVWCNGEEY RGAVDRTESGRECQRWDLQHP HQHPFEPGKFLDQGLDDNYCR NPDGSERPWCYTTDPQIEREFC DLPRCGSEAQPRQEATSVSCFR GKGEGYRGTANTTTAGVPCQR WDAQIPHQHRFTPEKYAC/KVR WAGGRALGRAAAGETGGG*SR A*GWRLAG*G*VERLLRDLREN FCWNLDGSEAPWCFTLRPGTR VGFCYQIRRCTDDVRPQDCYH GAGEQYRG\TV\SKTRKGVQCQ RWSAETPHKLQ*VPGAPGPAR ALTLGRHALMSGTRAWKWLR LSLPRFRSRSRLGSRLTSEPHAQ LEENFC/RDPDGDSHGPWCYTM DPRTPFDYCALRRCADDQPPSIL DPPPDQVQFEKCGKRVDRLDQ RRSKLRVAGGHPGNSPWTVSL RNR*GTTACLPQRGAEVVSSVV MPLGAGNLSLPPEVLARRWQV *HLSQESVPCPNSPLL*\QGQHF CGGSLVKEQWILTARQCFSS\CE PPLCLGTQSHPTFPFPQAS*QVS LGATD*ESQMT*QSFSPSHMPL
						TGYEVWLGTLFQNPQHGEPGL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hođ	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19619	49987	A	19734	727	3012	  PAPTHAHSAGQRSPLNDFQVLR
19019	49967	<b> </b> ^	19734	1,2,	3012	, , ,
						GTELQHLLHAVVPGPWQEDVA
						DAEECAGRCGPLMDCRAFHYN
	]					VSSHGCQLLPWTQHSPHTRLRR
						SGRCDLFQKKDYVRTCIMNNG
						VGYRGTMATTVGGLPCQAWS
						HKFPNDHKYTPTLRNGLEENFC
		l				RNPDGDPGGPWCYTTDPAVRF
	ŀ					QSCGIKSCREAACVWCNGEEY
						RGAVDRTESGRECQRWDLQHP
						HQHPFEPGKFLDQGLDDNYCR
						NPDGSERPWCYTTDPQIEREFC
						DLPRCGSEAQPRQEATTVSCFR
						GKGEGYRGTANTTTAGVPCQR
						WDAQIPHQHRFTPEKYACKDL
						RENFCRNPDGSEAPWCFTLRPG
						MRAAFCYQIRRCTDDVRPQDC
				]		YHGAGEQYRGTVSKTRKGVQC
						QRWSAETPHKPQFTFTSEPHAQ
				Ī		LEENFCRNPDGDSHGPWCYTM
						DPRTPFDYCALRRCADDQPPSIL
						DPP\DQVQFEKCGKRVDRLDQR
i						CSKLRVAGGHPGNSPWTVSLR
						l .
						NWQGQHF\CGGSLVKEQWILT\
	ŀ					AR\QCFSS\CHMPL\TGY*GMVG
						ATFFQNPQHGEPSL\QRVQ*AK
						MLCGPSGSQL\VLLKLERSVTL
						N/HRVWALIILP\PEWYVVPPGT\
						KCEIAGWGET\KGTG\NDTV\LN
				1		VALLNVISN\QECNIKHRGRVRE
						SEM\CTEG\LLAPVGACESDYG
						GP\LACFT\HNCWV\LEGIIIPN\R
19620	49988	Α	19735	66	234	WCSAWHSWLPHSPSGPAASYP
						PAACRSAPQWPDP*R*CPRTHR
İ						RRSPWRAPWHCP
19621	49989	Α	19736	3	500	GRGGGGVRRCARPAPGRPPAA
						ARRRGDQRGGRLGR\HVPGKTF
						CCGICGRGFGRRETLKRHERIH
						TGEKPHQCPVCGKRFRESFHLS
						KHHVVHTRERPYKCELCGKVF
						GYPQSLTRHRQVHRLQLPCAL
				İ		AGAAGLPSTQGTPGACGPGAS
	L	L			L	GTSAGPTDGLSYACSD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19622	49990	Α	19737	899	2348	GRRRGGGGRCWRGCQSGPERF
' ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' '						SCATCGQSFKHSW/DLVTHKYV
						HLMLLLPPSCEAVAPRGWRRSC
						SCPECCRVC/ELSSAA/RSATAR
						CFRRRSPSHAQPAPGPQVPQDW
		}				RSPRARIRLPVGPVGMPAFPTSA
		1				SGKAWPSLRSVSPHAFWKYHT
					1	SPPAMQPSGPLWASVPACAP*K
ŀ		ĺ				NIPSPTIWQRLAFSGLGLPICWV
		ł				ENHGPATSSQWLVSGPRSPHLL
						S/RAEHPFSRPLPRWVATAERQ
		1				RRPSGKRTNPFPASPAPPPGA/G
						PEVSVPPAGALPGEALGPARDL
			}			WPCSSWT*AGLPALG/DTGGGE
						APAP/AAAAEPSEDTLYQCDCG/
1						TFFASAAALASHLEAHSGPATY
		•				GCGHCGALYAALAALEEHRRV
		ŀ				SHGEGGGEEAATAAREREPAS
		ŀ				GEPPSGSGRGKKIFGCSECEKLF
						RSPRDLERHVLVHTGEKPFPCL
						ECGKFFRHECYLKRHRLLHGTE
						RPFPCHICGKGFITLSNLSRHLK
						LHRGMD
19623	49991	Α	19738	2	358	
19624	49992	A	19739	635	2441	LKAIKRHPFFVTIDWNVSCMPQ
						SPPPFKPAVGRPEDTFHFDPEFT
						ARTPTGVCHSGSLYNAHHLFR
						GFSFVASSLIQEPSQQDLHKVPV
						HPIVQVTVFACSGFTDGYEIKE
						DIGVGSYSVCKRCVHKATDTE
						YAVKVRQAC*VSSLVFPAGTG
		ŀ				APGIGEYGSGKLVSILCAQDSP
						HNLLPQRKVEKPSLSILVGESRS
						HVDKGQEVDILEMGMEFFFFIE
						FLVSPLATG*CFPMARSPCMGC
						SLSVCTAL*GQPLQGNTPTFKY
		1				QQVTWPQAKPPTPTAVLSPETA
		1				NVTSPQEQTRCLLKDRWLYPV
		1				VRNSSTRGPSEEIEILLRYGQHP
		1				NIITLKDVSGPRATASLVMELM
						RGGELLDRILRQRYFSEREASD
						VLCTITKTMDYLHSQGVGAMP
						GAAFNILYRDESGSPESIRVCDF
				1		GFAKQLRAGNGLLMTPCYTAN
						FVAPEVSIPRLQTQACDIWSLGI
						LLYTMLAG*AVTSNRCHDTPEE
						ILARIGSGKYALSGGNWDSISD
						AAKVSTCPSVALDPHQRLTAM
		ł				QVLKHPWVVNREYLSPNQLSR
	}	1				QDVHLVKVPAG\AVRTEWWE
L	<u>L</u>					WDLNRTPQAPRLEPVLSSNLAQ

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19625	49993	A	19740	144	1143	REAGRGCSAGVGGRPRRRRK GAASRARLPFSLSIMDPSLLRER ELFKKRALSTPVVEKRSASSE\S SSSS\SKKKKTKVE\HGGSSGSK QNSDHSNGSFNLKA\LSGSSG\Y KFGVLAK\IVNYMKDTGIQRGD THPLTLDEI\LDETQHLDIGLQQ KQWLMTE\ALVNNPKIEVIDGK YAFKPKYNVRDKKALLML\LD QHDQRGLGGILLEDIEEALPNS QK\AVKAL\GDQIPIC*SGPVRR KILF\F\NDKSCQFSVG*RIFRKL WEECQLVDSHGTEEKI\EEYLE ADRVFSSHARESG\PK\KVGPYS RRKGKSPAFTRKKPTFLRTSLT NHLAWNC
19626	49994	A	19741	677	1590	AHPHSPRRTLERQARISGAGRR SSPNAANRGSAPPSRRLASFCT AGSA\GRPA*SSPCSGPWTAVSA PRSPGPRGGGRPPSPGRRRPCCS VFSGGASPSSARVSSPSGSCSAS PPAAPAAWSLRGCGCWPG/VW EVHVNTGGDSV\AAPAAWGSR AACGPNLGCGSHPRAGRRAGH RCGHPWSPRPAPRPSCCGRSPR PGPPGASGCPRPAPRPPTPASRS GRPCSWRRARAAGWPRGWWG SSAGGCRRGSGRREWCPCGTA RGSPAAGVSRWTAPRCRHRRS DASASGTPARFPGAGRKGRTL WQ

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
NO:	sequence	nou	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	'			sequence		
10.60	1,000.5		1.05.15	la a	0.170	OVA (OD LEA WAREL CHOCK) THE
19627	49995	Α	19742	754	2179	SKMSRLEAKKPSLCKSEPLTTE
						RVRTTLSVLKRIVTSCYGPSGR
						LKQLHNGFGGYVCTTSQSSALL
						SHLLVTHPILKILTASIQNHVSSF
						SDCGLFTAILCCNLIENVQRLGL
						TPTTVIRLNKHLLSLCISYLKSE
						TCGCRIPVDFSSTQILLCLVRSIL TSKPACMLTRKETEHVSALILR
						AFLLTIPENAEGHIILGKSLIVPL
						1
						KGQRVIDSTVLPGILIEMSEVQL MRLLPIKKSTALKVALFCTTLS
					:	!
-						GDTS*\PGEGTVVVSYGVSLEN
		l				AVLDQLLNLGRQLISDHVDLVL
						CQKVIHPSLKQFLNMHP\IIAIDR
						IGSDSD/VEPLDLKWTGTQPIGS
				}		LGSICPNSYGSVKDVCTAKFGS
						KHFFHLIPNEATICSLLLCNRND
						TAWDELKLTCQTALHVLQLTL
						KEPWALLGGGCTETHLAAYIR
						HKTHNDPESILKDDECTQTELQ
10600	40006	<u> </u>	10742		652	LIAEAFCSALESVVGSLEHDGG LMPKPGRDTTKKENFRPISLMN
19628	49996	Α	19743	1	032	
				}		INAKILNKILANQIQQHIKKLIN HDQVGFIPGMQGWFNIHKSINV
						IHHIKRTNDKNHMIISIDAGKAF
						NKIQQCFMLKTLNKLNIHGVYL
ĺ	i					KIIRAIYD/KPPANIILNGEKLEA
İ						FPLKTGIRQG/CPLLRNIV/LEVL
						ARAIRQEKEIKGIQSGNEEVKLS
						LFADDMIVYLENPIVSAQNLLK
						LISNFSKVSGYKISVQ
10620	49997	1	19744	1	834	MGDFNTPLSTLDRSTRQKVNK
19629	49997	A	19/44	1	034	DIQELNSALYQVDLIDIYRTLHP
						KSTEYTFFSAPHHTYSKIDHIVG
		1				SKALLSKCKRTEIITNCLSDHSA
						IKLELTIKKLTQNRSTTWKLNN
		1				LLLNDYWKYKQPSENKHLYAN
						KLENLEEMDKFLDTYTLPRLNQ
		}				EEVESLNRPITRSEIEAITNSLPT/
		1				KKSPGPDGFTAEFYQ\MLEVLA
		1				RAIRQEKEIKGIQLVKEEVKLSL
						FADDMIVYLENPIVSAQNLLKLI
						GNFSKVSGYKI/NVQKSQAFLY
1						TNNRQTESQIM
19630	49998	В	19745	i	1521	LIMING I ESCHAI
19631	49998	В	19746	1	1461	
17031	72777	10	12740	<u> </u>	11701	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19632	50000	A	19747	1	2347	MELKTKARELHDECTSLSSRFD
		ĺ				QLEERVSVMEDEMNEMNLPTK
		1				KSPGPDGFTAEFYQRYKEELVP
						FLLKLFQSIEKEGILPNSFYEPSII
		l				LIAKPGRDTTKKENFRPISLMNI
						NAKILNKMLANQIQQHIKKLIH
						HDQVGFIPGMQGWFNIRKSINV
						IQHINRTKDKNHMIISIDAEKAF
	1	l				DKIQQHFMLKTLNKLVLEVLA
		l				RAIRQEKEIKGIQLGKEEVKVSL
						FADDMIVYLENPTVSAQNLLKL
						IGNFSKVSGYKINVQKSQAFLY
		1				TNNRQTERQIMSELPFTIASKRI
	İ	]		1		KYLGIQLTRDVKDLFKENNKPL
				ļ		LKEVKEDTNEWKNIPCSWVGRI
						NIVKMAILPKVIYRFNAIPIKLP
		l				MTFFTELEKTTLKFIWNQKRAC
						IAKSIFSQKNKAGGITLPDFKLY
	:					YKATVTKTAWYWYQNRDIAQ
	:					WNRTEPSEIMLHIYNYLIFDKPE
	1					KNKQWGKDSLFNKWCWENWL
						AICRKVKLDPFLTPYTKMNSR
						WIKDLNVRPKTIKTLEENLGITI
						QDIGVGKDFMSKTPKAMATKA
						KIDKWDLIKLKSFCTAKETTIRV
						NRQPTTWEKIFATYSSDKGLISR
						IYNELKQIYKKKTNNPIKKWAK
						DVNRHFSKEDIYAAKKHMKKC
						SSSLAIREMQIKTTMRYHLTPV
						RMAIIKKSGNNRKIQ/GGIWCD
						RIL*R*TTCRVAKEIQSL*RRI/W
	<u> </u>					KRLQRTLSIPVLDAV*PPMF*AS
19633	50001	Α	19748	1	1818	

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19634	50002	A	19749	I	1725	MELKTKARELREECRSLRSRCD QLEERPNLRLIGVPESDGENGT KLENTLQDIIQENFPNLAKQVN VQIQEIQRTPQRYSLRRATPRHII VRFTKVEMKEKMLRAAREKGR VTLKGKPIRLTADLSAETLQAR REWGPIFNILKEKNFQPRISYPA KLSFISEGEIKSFTDKQMLRDFV TTRPALQELLKEALNMERNNRS WFFEKINKIDRPLARVIKKKRE KNQIDAIKNDKEDITTNPTEIQTI IREYYKHLYPNKLENLEEMDTF LDTYTFPRLNQEEVESLNTSITG SEIVAIIS/NSIPTKKSPGPDGSTA EFYQMLEVLARAIRQEKEIKGI QLGKEEVKLSLFADDMIVYLEN PIVSAQNLLKLISNFSKVSGYKI NVQKSQAFLYTNNRQTESQIMS ELPFTIASKRIKCLGIQLTRDVK
						DLFKENYKPLLKEIKEDTNKW KNIPCSWVGRINIVKMAILPKVI YRFNAIPIKLPMTFFTDLEKTTL NFIWNQKRARITKSILSQKNKA GGITLPDFKLYYTATVTKTAW YWYQN\RWYWYQNRDIDQWN RTEPSEITPHVYNYLIF

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19635	50003	Α	19750	1	1719	MHNTDGNRFLSHWGYRQALSI
		1				SKPASASLHPSSKTKPLGTQSKT
						VVAKRNREHGKKERSSSPAME
		1				QSWMENDFDELREEGFRRSNY
		ł				SELREDIQTKGKEVENFEKNLE
		]				ECITRITNTEKCLKELMELKTK
		}		<u> </u>		ARELREECRSLRSRCDQLEERR
				:		KQERSKIDTLTSQLKELEKQEQ
			ļ			THSKAGRRQEITKIRAELKEIET
						QKTLQKINESRSWFFERINKIDR
						PLARLIKKKREKNQIDTIKNDK
			}			GDITTDPTEIQTTIREYYKHLYA
						NKLENLEEMDTFLDTYTLPRLN
		İ				QEEVESLNRPITGSEIVAIINSLP
	1					TKKSPGPDGFTAEFYQRYKEEL
						HINRAKDKNHMIISIDAEKAFD
						KIQQPFMLKTLNKLVLEVLARA
						IRQEKEIKGIQLGKEEVKLSLFA
						DDMIVYLENPIVSAQNLLKLLS
Ì	1					NFSKVSGYKINVQKSQAFLYTN
İ	İ					NRQTESQIMSELPFTIASKRIKY
						LGIQLTRDVKHLFKENYKPLLK
						EIKEDTNKWKNIPCSWV\GRINI
						VKMAILPKVIYRFNAVPIKLPM
					:	TFFTELEKTTLKFIWNQKRACIA
						KSILSQKNKAGGITLP

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19636	50004	A	19751		2133	WRRIYQANGK*KNK/QKKAGV VILVSDKTDFKPTKIKRDKEGH YIMVKGSIQQEELTVLNIYAPN TGAPRFMKQVLRDLQRDLDPH TTIMGDFNTPLSTLDRSARQKV NKDIQELNSALHQADLINIYRIL HPKSTEYTFISAPHRTYSKIDHI VGRKALLRKYKRTEIITDCLSD HSAIKLELRIKKLTQNSSTTWK LNNLLNDYWIHNKTKAEIKM CFETSENKDTTYQNLWDTCKA VCREKFIALNAHKRKQERSKID TLTSQLKE/LEKQEQTHSKASRR KSRRNG*IPGHIHPPKTKPGRI* VPE*TNNRV*N*GNN**LTNQK KFRTRIHSQILPEHSAGSSGQG NQAGERNKGYSIRKRGSQIVPV CR*HDCIFRKPHHLSPKSP*AVK QLQQSLRIQNQRAKITSSPIHQ* QTNREPNHE*TFIHNCFKENKIP RNPTYKGCEGPIQGELQTTAQQ NKRGHKQMEEHSMLMDRKNQ YHENGHSAQGNL*IQCHPHQAT NDFLHRIGKNYFKVHMEPKKSP HCQVNPKPKEQSWRHHAT*LQ TILQGYSNQNSMVLVPKQTYRP MEKNRGLRNNTTHLRPSSL*QT *QKQEMGKGFPI**MVLGKLAS
						HM*KAETGSLPYTLYKN*FKM D*RLKC*T*NHKNLRRKPRQYH SGHRHEQGLYV*NTKSNGNKS QN*QMGSN*TKELLHSKRNYH
19637	50005	Α	19752	1	1314	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19638	50006	A	19753	1	1998	MVKGSIQQEELTILHIYAPNTG
17030	30000	l'`	17733	1	1990	APRFIKQVLSDLQRDLDSHTLI
						MGDFNTPLSTLDRSMRQKVNK
						DTQELNSALHQAALIDIYRTFH
						PKSTEYTFFSAPHHTYSKIDHIV
						GSKALLNKCKRTEIITNYLSDHS
						AIKLELRIKNLTQSRSTTWKLN
						NLLLNDYWVHNEMKAEIKMFF
		į				
		į				ETNENKDTTYQNLWDAFKA\PL
		į				ARLIKKKREKNQID/TIKNDKGD
						ITTDPTEIQTTIREYYKHLYANK
[						LENLEEMDTFLDTYTLPRLNQE
ĺ						EVESLNRPITGSEIVAIINSLPTK
						KSPGPDGFTAEFYQRYKEELVP
						FLLKLFQSIEKEGILPNSFYEASII
						LIPKPGRDTTKKENFRPISLMNI
					:	DAKILNKILANRIQQHIKKLIHH
	İ	:				DQVGFIPGMQGWFNIRKSINVI
						QHINRAKDKNHMIISIDAEKAF
						DKIQQPFMLKTLNKLVLEVLAR
						AIRQEKDIKGIQLGKEEVKLSLF
						ADDMIVYLENPIVSAQNLLKLIS
						KFSKVSGYKIHVLKSQAFLYTN
						NRQTESQIMRELPFTIASKRIKY
						LGIQLTRDVKDLFKENYKPLLN
						EIKEDTNKWKNIPCSWIGRINIM
						RRSSSAPSGCHLCPPSLYLHSPN
						VELKQASFSVAGFQWKLGITAP
	1					TQPRGVGDDFVTSTSPPSLDMF
						PRLKGAENSPHRSGRD
19639	50007	Α	19754	1	3288	
19640	50008	A	19755	1264	1293	LUZZI DZI ANI VEDDE OTEGODIC
19641	50009	A	19756	264	490	VYLLIVLAVLYTNNRQTESQIM
		İ				SELPFTIASKRIKYLGIQL\TRDV
		l				KDLFKDNYIPL\LKEI*EDTSKW
10640	50010	ļ. —	10767	1	1045	KSIPCSWI
19642	50010	A	19757	1	1845	MCDENTEDICTI DECEDOMANIC
19643	50011	Α	19758	1	3144	MGDFNTPLSTLDRSSRQKVNK
						DTQELNSTLHHADLIDIYRTLHP
					}	KSTEYTFFSAPHHTYSKIDHVV
		l				GSKALLSKCKRTEIITNCLSDHS
		l				AIKPELRIKKLTQNRSTTWKLN
		1				NLLLNDYWVHNKMKAEIKMFF
						ETNENKDTTYQNLWDTFKAVS
						RGKFIALNAHKRKQKRCKIDTL
						ASQLKEVEKQEQTHSKASRRQ
						EITKIRAELKEIETQKTLQKINES
	1	I	Į.	I		RSWFLERINKIDRPLARLIKKKR
	ł	1				
19644	50012	В	19759		1743	EKNQIDVIKNDK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19645	50013	A	19760	<u> </u>	1500	MGKKQSRKTENSKNQSASPPPE
						ECSSSPAMEQSWMENDFHELR
						EEGFRSNFSELKEDVRTHSKEA
						KNLEKRLDEWLTGITSVEKSLN
						DLMELKTMARELCDKCTSFSS
			•			KFHQLEERVSVTEDQMNEMKS
		İ				WFFEKINKIDRLLARLIKKKRE
						KNQIEAIKNDKGDITTDPTEIQT
						TIREYYKHLYTNKLENLEEMDT
						FTGTYTLPRLNQEEIESLNKPIT
						GSEIEAIINSLPTKKSPGPDGFTA
						EFYQRYKEELL\FNIRKLINLIQL
						INRTQDP*PPL/VLSRIGRRVHQ
						YLENPIVSAQNLLKLKSNFSKV
						SGYKINVQKSQAFLYIKKRQTE
						SQIMSELPFTIASKRIKYLGIQLI
		l		i		RDVKDLFKENYKPLLNEIKEGT
						NKWKNIPCSWIGRINIVKMAILP
		l				KVIYRFNAIPIKLPMTFFTELEKI
						TLKFIWNQERACIAKTILSKKN
						KAGGIMLPDFKLYYKA\TVIKT
		1				AWDWYQNRYLDQWNRTEPSEI
		<u> </u>				TPHIYNHLIF
19646	50014	Α	19761	1	3192	
19647	50015	В	19762	180	190	
19648	50016	Α	19763	1	3212	MVKGSIQQEELTILNIYAPNTG
						ALRFIKQVLRDLQRDLDSHTIIM
					-	GDFHTPLSTLDRSTRQKVNKDI
1						QELNSALHQEDLIDIYRTLHPKS
						TEYTFFSAPHHTYSKIDHIVGSK
		l				ALLSKCKRTEIITNCLSDHSAIK
						LELRIKNLTQNRSTTWKLNNLL
						LNDYWVHNEMKAEIKMFFETN
						ENKDTTYQNLWDTFKAVCRGK
		1				FIALNAHKRKQERSKIDTLTSQL
						KELEKQEQTHSKASRRQEITKIR
	L		]			AELKEIETQ

SEQ ID NO:	SEQ ID NO: of peptide sequence	ı	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	l e	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide inscrtion)
19649	50017	A	19764	1	2607	MTNHCFKRQFQKHEMLLKEDC EPGDSSFILGEQHWENMLCEPK QHPPQQGEGRAAGQAQNPLNT
						DVSPGAGAKESEEKIYQHLNYT YFKQKCMSCSGFTFSSYAMHW
						VHQAPGKGLEWVSAIGAGGGT
						YYADSVKGRFTISRDNAKNSLY
ļ.						LQMNSLRAEDMAVYYCARDTI
ŀ						DAYGDEKPSTIVTRIRVLRKLR
		İ				DTCHMGEADRTFLSPIAQHPPV
						CHVETHLGMQVANSSEEDELV
						WSKWDVDVKFILCCAKYHREQ
						LQESALDLKSLHCPSASPALFL
						ATPSQPVLTTGPGSASSQGGAE
						MGAHFLVQLVQSGAETRESFD
						PALPEMETQRTDEQCRAVHPGP
						QKERHGNMSPPLLMKGSSSPVP
						SGPGEEPSHREPVSFICGRCILLT
		1				LLLGDCLWGEGAWTQGDVLQ
						PSDRASFLAMGVNTTVQQVGD
	}					LSGYFPNSVGKACKCREFHTLT PLAHTSSTTHETFPGAQCEVQL
						VESGGGLVQPGGSLRLSCPDSG
						FTFSNHYMSWVRQAPGKGLEW
						ISYISGDSGYTNYADSVKGRFTI
						SRDNANNSPYLQMNSLRAEDT
						AVYYCVKHTARGKLEKQEQTY
}						SKASRRQEITKIRAELKETETQK
						TLQKINESRSWFFEKIYKIDRPL
						ARLIKKKREKNQIDAIKNGKGD
						ITTDPTEIQTTLREYYKHLYAN
						KLENLEGMDKFLDAYTLPRLN
19650	50018	Α	19765	1	1251	
19651	50019	Α	19766	1099	4113	
19652	50020	A	19767	1	2628	
19653	50021	В	19768	1	2970	
19654	50022	В	19769	1	4142	
19655	50023	Α	19770	1	4091	MHIVVETALSASWQNKAKPPA
						RVLLQVVPNVWFLVAVVWEL
						YPSLDLMDRSIECSSSPATEQS
						WTENDYDKLREEGFRRSNYSE
						LQEDIQTKGKDVENFEKNLEEC
						ITRITNTQKCLKELMELKTKAR
						ELREECRSLRSRCDQLEERVSV MEDEMNEMKQEGKFREKRIKR
						NEQSLQEIWDYVKRPNLRLIGV
						PESDGENGTKLKNTLQDIIQENF
						PNLARQAKVQIQEIQRMPQRYS
						LTRATPRHIIVRFTKVE
	L	L	L	L	l	LIKATIKIIIVKETKVE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19656	50024	Α	19771	156	450	LQVEIYKTVKMHCIWHKQMVS
						SYFFYQKKNELSNFEEVFLKTA
						SGMFFSTVGCHPTRCGEFEKNN
						PDLYLKELLNLAENNKGKVV\A
						IGECGLDFDRLQ
19657	50025	Α	19772	173	1009	LHGGNLQDSKDALHLAQTNG
		ŀ				MFFSTVG\IHPTRCG*/YFEKNNP
						DLYLKGVAKIFA/EKTLKGKLW
						QLGEC\GLDFD\RLQFCPKDTQL
		l				KYFEKQFE\LSEQTKLPMF\LHC
						RNSHGWNFLDIMKRNRDRCVG
		1				GVVHSFDGTKEAAS\SLIDLDLY
						IGFNG/CASPENLEANLGRFWK
		İ		1		SI\PSEKLMIEA\DAP\WCGVKST HA\GSKYIRTAFPS\KKKWESGH
		l				CLKDRNEPCHIIQILEIMSAVRD
						EDPLELANTLYNNTIKVFFSWN
		İ				IIGICLPLSIMYVKFHSKTS
19658	50026	A	19773	16	1558	QEHLQRASPRLLQLLPPLPPPPP
17030	30020	l' <b>`</b>	17773	1.0		PPAFGGSLSSLGSRMSRQV\VRS
		ŀ				SKFRHVFGQPAKADQ\CYEDVR
		}				VSQTT\WDSGFWC/SVNP*VLW
						ALICEASGGGAFLVLPLGKIWT
						LWDKNAPTGCGSHS/APVLDIA
						WCPHNDNVIA\SGSED\CTVMV
		İ				WEIPDGGLMLPLREPVV\TLEG
		ł				HTKRVG\IVAWHTTAQNVLLSA
			<u> </u>			GCDNVIMVWDVGP\GAAMLDT
						GPRGA/HPDANYRGDWEP\DGG
						LICTSCRDKRVRIIEPRKGTVVA
1						EKDRPHEGTRPVRAVFVSEGKI
						LTTGFSRMSERQVALWDTKHL
						EESPV/SLRKLHTRNGVPVAFFG
						P\DTNIVYLRGKG\SGPIRVFE\IT
						SE\APFLHYL\SMFS\SKESQRG\
						MGYM\PKRGRKVNK\CEIAGFY
						KLHDGR/CVSPMPKPVPRKSDL
						FQEDLGPTPPRGPDP\ALTA\EE
						WLGGRDVG\PLFISLKDGYV\PP
						KSREL\RVNRGLDP\GRRR\AAP
						EASGTPSSDAVSRLEEEMRKLQ
10650	60027	<del> </del>	10774	220	1475	ATVQELQKRLDRLEETVQAKL
19659	50027	Α	19774	229	475	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19660	50028	Α	19775	225	965	EWAWRSQLRSEHSRADPTRSSC
						RSCGAKMSGEENPASKPTPVQ
			ļ			DVQGDGRWMSLHHRFVADSK\
						DKSGSKKKNPPPPEVVFIGDSL
	İ					VQLMHQCEIWRELFSPLHAL\N
						FGIGGDGTQHVLWRLENGELE
						HIRP\KIVVVWVG/TPNNHGHTA
						EQGLLPRGQHPNPLREKNRQV
						NELGTGGTWLGHPRAHFLDAD
	1					PGFVHSDGTISHHDMYDYLHLS
						RLGYTPVCRALHSLLLRLLAQD
		1				QGQGAPLLEPAP
19661	50029	В	19776	691	1077	
19662	50030	Α	19777	481	1308	PSVAFL*PSLHRYGHTGTAGPG
						WPCAAGS*AA/SSSRRGLPSVLQ
						ASSNSSHRRCNSTPSGSAGQRR
						VRRAP*MASTTPGGTESGPCKA
						RNAFRMRSTSAFCRGQ*QRPVP
						SSSSRSSSRACA\AGETASGDP
ŀ		1				RAQRQLRS*DEESSSPSLDFRQA
						/PWPTAPCVKVSYSSSRMFSGM
						SPWRSSDNKRNSSFLRKTTGSN
						FSADLDTPPMPEVCLLPSLGYP
						LVPPVAEVHMPVPAPILPSAGIR
						RKIQKAASAKQLPLFQMDDSFS
						QHRVQGDTKK
19663	50031	С	19778	41	318	
19664	50032	Α	19779	970	1193	INNHIRDSCWTCHCSC/QPPPPPP
1						LLPQPPP*LSAPRVIERKE*DRL/
İ						CPGEQTSSPTPSPRTQSAAAAT
L						QLSLPPP
19665	50033	Α	19780 .	1	494	MDGRVMAAGGRQAPGQKKAG
						TSEAPPSSQEWPEAWYGCLQQ
						KLRAVYLVQSQPCTGPVLAPG
						AACPAAAASTPGCAQW/CGPR
						ACSPTQSSQLCAWFAFSSRCPD
						FCPNPPRPL*DPPSWPTDPADRC
						SPELTIRDSCWTCHCSCRRRHR
		L				RPCCRSRRPDSPRHG
19666	50034	C	19781	120	392	
19667	50035	Α	19782	174	375	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19668	50036	A	19783	33	1154	QFALCAGWWLCGRGGGRTPAT MSGFSTEERAVPFSLEYRVFLK NEKGQYISPFHDIPIYADKDVFH M\VVEVPRWSNAKMEIATKDP LNPIKQDVKKGKLRYVA\NLFP\ YKGYIW\NYG\AFPQTW\EDPG\ HNDKTLLGCCGLTMDP\IDVC* NWEGKVCARGEIIG\VKVLG\IL AMI\DGGGTDWKVIAINV\DDP GCQPINNDI\NDVKRLK\PGYL* KLPVDWFR\RYKVPDGKPENEF AFNAEFKDK\DFA\IDIIKSTPWT LGKALVT*ENRMGKGISCMNT NFVLRAPFKVCILDAAQSPLWD ALPTHPCGICPCHSTQQTVDKG VPITQKNLMRFPLEYQADIAYIR VPSWMVLEVKVVAFSKAFKFV
19669	50037	A	19784	1	612	EPHPNLK  VTQEDDIIEEHKPRTPRPSPPLL  APSPACQAFGEGHSPGHPAIST  GAGPRPSASFSHPHLSGLHREPP  LNWSVCPGLVWSPKPALDETH  PFVQQFPHQATLEWGVFHRDV  GRRWGRRLGNLQIGSEVARRR  RVGGGGLQCRRLGVGASPGAD  W**SG*EEACGGRWAA/GADG  WGLGRPRELIGCRGWRRSLAA  APGPQRPGFVSVSTPRCRPRSG  METQTPALRVQQDDLFQS
19670	50038	A	19785		951	MEEGISSQIGRRENECQQKKCQ TLIKPSDLVRTHSYHENSMGETI LVIQLPPPGTAFDMWELQFKLI TQTFSHHNQLAQKTRREKRAR QEA\ERREKAERAARLAKEAKS ETSGPQIKELT\DEEAERLQLEID QKKDAENHEA\QLKNG\SLDSP GKQDTEEDEEEDEKDKGKLKP NLGNGEDLPNYRWTQTLSELD LAVPFCVNFRLKGKDMVVDIQ RRHLRVGLKGQPAII\DGELYNE VKVEESSWLIEDGKVVTVHLE KRGPADAHSRALSVLQASRLL GVHCVPGTVPNAEIQVVDFVIY FWQSVIPQASPISE

Possible nucleotide ble nucleotide insertion)  RSAARLGGEQEEER IAQ\QHEGGVQELV KTDFFIGGEEGTAE IHNQL\AQKT\RREK REKAERAARLAK PQIKELTDEKA\ERL DAENHEAQLKNGS FEEDEEEEDEKDKG RADLAQ*RLNPDPC
RSAARLGGEQEEER IAQ\QHEGGVQELV KTDFFIGGEEGTAE IHNQL\AQKT\RREK RREKAERAARLAK PQIKELTDEKA\ERL DAENHEAQLKNGS IEEDEEEEDEKDKG RADLAQ*RLNPDPC
IAQ\QHEGGVQELV KTDFFIGGEEGTAE IHNQL\AQKT\RREK RREKAERAARLAK QIKELTDEKA\ERL DAENHEAQLKNGS IEEDEEEEDEKDKG RADLAQ*RLNPDPC
IAQ\QHEGGVQELV KTDFFIGGEEGTAE IHNQL\AQKT\RREK RREKAERAARLAK QIKELTDEKA\ERL DAENHEAQLKNGS IEEDEEEEDEKDKG RADLAQ*RLNPDPC
KTDFFIGGEEGTAE IHNQL\AQKT\RREK RREKAERAARLAK PQIKELTDEKA\ERL DAENHEAQLKNGS IEEDEEEEDEKDKG RADLAQ*RLNPDPC
IHNQL\AQKT\RREK RREKAERAARLAK PQIKELTDEKA\ERL DAENHEAQLKNGS FEEDEEEEDEKDKG RADLAQ*RLNPDPC
RREKAERAARLAK PQIKELTDEKA\ERL DAENHEAQLKNGS FEEDEEEEDEKDKG RADLAQ*RLNPDPC
QIKELTDEKA\ERL DAENHEAQLKNGS FEEDEEEEDEKDKG RADLAQ*RLNPDPC
DAENHEAQLKNGS FEEDEEEEDEKDKG RADLAQ*RLNPDPC
TEEDEEEEDEKDKG RADLAQ*RLNPDPC
RADLAQ*RLNPDPC
'
" (" DIED! " (" )
F\CVNFRLKGKDMV
\VGLKGQPSDSLM
VEES\SWLIEDG\KV
NKMEWWSRLVS\S
INPENSKA\SDL\DS
KDDGMTQRTESPM
RKQEILK\KFM\DQ
KAKFPTNPCFFPP
VAMAPTIQTQAQ
SHRTLPERSGVVCR
DIPFDPKFITYPFDQ
TSLEKQHKHDLLT
DLINPDTYRIDPNVL
LEEEIQAPTSSKRSQ
V\MRKTEYISTEFN
PEVKKWGSVKQQF
DSQITAIEKTFEDA
WGEARISQHYSKP
PVFPDFKMWINPC
APKDTSGAAALEM
GMMDEEGN\QFVA
KKRKRDQEEEMD
YKIAREYNWNVK
ENYFFIFREGDGVY
RLSKRRAKA\GVQ
VKHRDMNEKELEA
ENHEPEEEEEEEME
SDEEQEKGSSSEKE
SESEREEGDRHEAS
DSSDY*ARAARDK
SEDDADSDDEDRG
DSDSGRN\GGGQR
PFPSGSEHSAQENG
SEADSDSD

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19677	50045	A	19792		1344	PGRTTRSMAEDVFLSAPIPRGC ADGRDADPTEEHMAQTERNDE EQFECQELL*CHVQVGAPEEEE EEEEDAVLVAEAEAVAAGWM LNFLCLSLCRAFREGRSEDFRRT RNSAEAIIHGLCSLTACQLRTIYI CQFLTRIAAGKTLDAQFENDER ITPLESALMIWGSIEKEHDKLHE EIQNLIKIQAIAVCMENGNFKG AEEVFERIFGDPNS\HMPFKSKL LMIISQKDTFHSFF\QHFSYNHM MEKIKSYV\NYVLSEKSSTFLM KAAAKVVESKRTRTITSQDKPS GNDVEMETEANLGYKKKC*LT NSLR*LNPQRVQYPY*GSHKNL FYLKLQHGTQQQDLNKKERRV GTPQSTKKKKESRRATESRIPVS KSQPVTPEKHRARKRQAWLWE EDKNLRSGVRKYGEGNWSKIL LHYKFNNRTSVMLKDRWRTM KKLKLISSDSED
19678	50046	Α	19793		1444	IPGSTISCLKGQYPSEPFNMAED VSSAAPSPRRCADGRDP\DPTEE QMAETERNDEEQFEC\QELLEC QVQVGAPEEEEEEEEEDAGLV AEAEAVAAGWMLDFLCLSLCR AFRDGRSEDFRRTRNSAKAIIY GLSMLTACPLRTIYICQ\FLTRIA AGKTP*MHRFENDE\RITPLGISP *MIWGSIEKEHDKLHEEIQNLIK IQAIAVCMENGNFKEAEEVFER IFGDPNSHMPFKSKLLMIISQKD TFHSFFQHFS\YNHMMG\KIKSY VNYVLSEKSSTFLM\KAAAKVV ESKRTRTITSQDKPSGNDVEME TEANLDTRKRSHKNLFLSKLQH GTQQQDLNKKERRVGTPQSTK/ RRKKEGRRATESRIPVSKSQP\V TPENDRARK\RQAWLWGRKDK ILRS\GVREFG\EGN\WS*ILLHY KF\NNR\TSVM\LKARW\RTMEG NLKLDFLQDSEDWIVFVKSFDG KGQFKYFDHCILFETCVIDVI
19679	50047	Α	19794	1	1020	
19680	50048	A	19795	411	769	SHLGFPGTWTRLWPSRMSPCPP QQSRNRVIQLSTSEL/APALGN WARLPQVTEQ*FRGLRKGSRT MPGSSPPATWTKGCLLA*Q*CL CPQLGWSRVAP*CRLGALFLAC SLLLAAPLRCF

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19681	50049	A	19796	171	325	ETVAVSSAGARASSIMRIMVSS RPSETSLESVFPDSAP/PGL*PPP CLSFGL
19682	50050	Α	19797	335	1536	GCSPCPPQ\QS\RNRVIQLSTSEL GEMELTW\Q\EIMSITEACRVLN APSEPSFEAPKPQLPYL\GPPPPT TY\CPC\SIHPDSGFPI\P\PPPYEL P\ASTSHVPDPPYSYG\NMAIPV SK\PLSLSGLLSEP\LQDPLALLD IGLPAGPPKPQEDPESDSGLSLN YSDAESLELEGTEAGRRRSELC RRCYPV\EYPYSLMPNSLAHSN YTL\PAAETPLALEPSSGP\VRA KPTARGE/AQGSRDERRAL\AM KISF\PTDKIVNFPVDDFNELLA RYPLTESQLALVRDIRRRGKNK VAAQNCRKRKLET\IVHLERKL ERLTNE\RERL\LRA\RGEASR\T LKVMRQQ\LQELY\REHFPSHLR E*NPGQTAYSP*KK*RACKQAC PNGDPSFPCGPRGT\KMEAHRL
19683	50051	Α	19798	I	408	LNLMRWEAVQARGGAFRIPRV DPGPTPIPHNSSSRFDCNSVSTA P\RVLVSGICAAGSFVLVAFSHS VGTSLC/GYFLLLTSPEAQDPGG EEEAESAARQPLIRTEAPESKPG SSSSLSLRERWTVFKGLLWYIV PLV
19684	50052	A	19799	1	759	
19685	50053	Α	19800	1	1815	·
19686	50054	A	19801		942	MGGCAGSRRRFSDSEGEETVPE PRLPLLDHQGAHWKNAVGFW LLGLCNNFSYVVMLSAAHDILS HKRTSGNQSHVDPGPTPIPHNS SSRFDCNSVSTAP\RVLVSGICA AGSFVLVAFSHSVGTSLC/GYFL LLTSPEAQDPGGEEEAESAARQ PLIRTEAPESKPGSSSSLSLRER WTVFKGLLWYIVPLVVVYFAE YFINQGLVSEGCLFRPFPFSALN VCEVNLVKVSQHQGGWVAVV TQGKKHPISCRRLNVDGMNIIS NDGSNKLNRSRKARVSYKVQR TINASLNAATSTRAVSSNPNFV

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19687	50055	A	19802	82	1620	SLSPVADPHPSRGSPLWTLYDP GPSGDLNLMRWIGCAGSRRRFS
						DSEG\EETVPEPRLPLLDHEGAH
						WKNAEGFWLLGLCNNFSYVV
						MLSAAHDILSHKRTSG\NQSHV\
						DPGPNGDSPTTAYSRFDCNSVF\
						TGAVLL\ADILPTLVIKLL\APLG
						L\HLLPYRSGPRVLVSG\ICAGG
						KFRPGFAFSHSVG\TSLCGCWSF
						ASISSGLG\EVTFLSLTAFYPRA
		-				VISWWSSGTGGAGLLGALSYL
		-				GLTQAGLSPQQTLLSMLGIPAL
					:	LLASYFLLLTSPEAQ\DPERGEE
•						AK\SQAGQPFIRTEGPE\GKPGSS
						SSLS\LRERWTVVKGLLWVPLF
						PWFEVTFPKYFITQGLFELP/LFF
						WTTSLSHAQQYRWYQMLYQA
						GVFASRSSLRS\CSIGFTW\ALAL
		1				LQCL\NL\VFLLADVWFGFLPSI
						YLVF\LII\LYEGAPWGSASLREK
						PF\HNIRPWRPSDEHREVCKWR
						PPCHL*HTGGFILSGASLALPLH
						DFLLPSSPDTRDPQDAGHIHLW
						ARGTGYQ
19688	50056	С	19803	270	542	
19689	50057	A	19804	271	449	CLELRNLFYGACSWEVSISSDS EKPGIFNASPASAVHC**GFARP SSWRVLGAICTFHHSVYAKTA
						KRKAST*RHFHH
19690	50058	Α	19805	21	79	
19691	50059	Α	19806	1242	1560	WATRSSCSSMMEMPPWCPCGC
						AGRWRRGGASRGPREAAAGH*
						KPRTSNQGEQPTRRPSPYAACA
	į					RRAFAGPWASLPSPSPLSAGGL
	<u> </u>					AWGAPCVAHPTRPSPYLLV
19692	50060	В	19807	148	2949	
19693	50061	Α	19808	1335	2173	KQAHRPREEKRREEKRKEG\PS
						VPETLKKKQRNFTELKIKRL\RK
						KFAQQML\RKAR\RKLIYEKAK
						HYPKE\YRKMYRT*ISNGRGWA
						RK\AGQLYLVLKNPKFAV\VIRN
						QKVSMGVSP\KLRKVVVSFLRP
						SSKSSNETFVKALKQRPSD*TM
						P*GLVRGPI*LPWGYPPQS*SPV
						NEP*SYK\RGY\AKIQ*EAELPLT
1						DNALIASISWVNSAFICM\DDLI
						HEIYTV\GKRFK\EANNFLWPLQ
						IVLLPRGG\MKKKTTHFVEGGD
10.55 :	50075	<del>ا</del> .	10000		100	AGTQRSDQINRLIRRMN
19694	50062	A	19809	3	186	
19695	50063	Α	19810	272	428	<u> </u>

NO: Of peptide hod in USSN location of first	codon for last amino acid	
1 1 10000000000 1 1 1 1 1 1 1 1 1 1 1 1	l l	*=Stop codon, /=possible nucleotide
sequence 09/540,217 codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
sequence		
19696   50064   A   19811   50	2237	RSLKAFICSILSMDKQSRDPPVK
		MQGSITTPGSIALAQAQAQAQV
		PAKAPLAGQVSTMVTTSTTTTV
		AKTVTVTRPTGVSFKKDVPPSI
		NTTNIDTLLVATDQTERIVEPPE
	:	NIQEKIAFIFNNLSQSNMTQKEE
		LVNNTELVQSYRQQIGNVVNQ
		ANLQLFWNMYNSRRDLDINRP
		GTVPNAKTLRCPVMLVVGDNA
		PAEDGVVECNSKLDPTTTTFLK
		MADSGGLPQVTQPGKLTEAFK
		YFLQGMGYMPSASMTRLARSR
		TASLTSASSVDGSRPQACTHSE
		SSEGLGQLNLKFEIEVLCKNLA
		LDINELKPGNLLKDKDRLKNLD
		EQLSAPKKDVKQPEELPPITTTT
		TSTTPATNSTCTATVPPQPQYS
		YHDINVYSLAGLVPHITLNPTA
		HPQLKQCVRQAIERAVQELVHP
		VVDRSIKIAMTTCEQIVRKDFA
		LDSEESRMRIAAHHMMRNLTA
		GMAMITC/RE/PLLMSISTNLKN
		SFASALRTASPLQREMMDQAA
		AQI*AQDNCFLL/CKVGGVDPK
		QLAVYEEFARNVPGFLPTNDLS
		QPTGFLAQPMKQAWATDDVA
		QIYDKCITELEQHLHAIPPTLAM
		NPQAQALRSLLEVVVLSRNSRD
		AIAALGLLQKSMENGLNYMAV
		AFAMQLVKILLVDERSVAHVT
		EADLFHTIETLMRINAHSRGNA
		PEGMDITTRQKVNKKREGFND
19697 50065 A 19812 1	3777	KNPEFNKMVLNETYRNIKVLLT
		SDKAAANFSDRSLLKNLGHWL
		GMITLAKNKPILHTDLDVKSLL
		LEAYVKGQQELLYVVPFVAKV
		LESSIRSVVFRPPNPWTMAIMN
		VLAELHQEHDLKLNLKFEIEVL
		CKNLALDINELKPGNLLKDKDR
		LKNLDEQLSAPKKDVKQPEELP
		PITTTTTSTTPATNTTCTATVPP
		QPQYSYHDINVYSLAGLAPHIT
		LNPTIPLFQAHPQLKQCVRQAIE
		RAVQELVHPVVDR

SEQ ID	_	Met	SEQ ID NO:	Nucleotide		Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	t .	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19698	50066	Α	19813	160	1002	KRLGSSAISMSKYKLIMLR\HGE
				i		GAWNKENRFCSWVDQ\KLNSE
						GMEEARNCGKQLKALNFEFDL
						VFTSVLNRAIPTARL\IL\EELGQ
						EWVPVESSWRLNERH\YGAL\I
						GLNREQ\MA\LNHGE\EQVRL\W
						RRSYNVTP\PPIEE\SHPY\YQEIY
						NRPGGIKVCDVP\LDQLPRSES\
						LKDVL\ERFLPYW\NERIAPEV\L
						RG\KTILISAHGK*AVGALLKHL
						GRYPSGLKTSINIYSFLLGVPILL
						\ELD\ENLRAVGPH\QFLGDQEAI
						QAAIKKVEDQGKVKQAKK
19699	50067	Α	19814	200	1155	NHYISLFQYSQDTCKLLFRNQM
19099	30007	^	17014	200		TISQIFLCIFSGVLMVFFLIITTSQ
						VSFTNFFTGLKIFFLVIKFWVSE
						FLHHQTGTYIHIYLLFISMKSFF
						LYASPLTFMILSHFQASLSSIFGF
ļ						CSSLFRQSSFFSNILFF*VYTQSY
						REFRTYYETIPCQFDITFVIFKM
						CCILILC\FDLLNRRNGSIIIQKFG
ł						IYGLVKIFIF*IFTNIHLLQ*FFQV
1						VFILQFFNIVKKSFLQPAGSKR*
						KFFSKVRFFSFL*F*KWEARSSA
						SKLFLLIY**VYR/IFSFLLVFPV
						V/CPVGWYLLFFFFLYLFFLFLN
						LSKNSFYSQWYWFKR
19700	50068	A	19815	l	168	RPWLHLHGRPGPWPAETPPPQD
19700	30008	^	17013	'	108	PEQAPPPGRAGPAGEAEGLSCV
						/P*APSRRPRTM
19701	50069	A	19816	1	531	PVGQGARQDDSSSSASPSVRAT
17701	30007	^	17010	1		SQGGHLMSDLFNKLVMRRKGI
						SGKGPGAGEGPGGAFVRVSDSI
						PPLPPPQQPQAEEDEDDWES*G
						AP*HLPPPDPDLGRCSDMDTAR
			į			TSCSDLLPWEGVTEPAVCGDQL
						QGAEGWLEATQLGRGLVCLYF
						MGGRLHPACATVFLRGFQRKR
						HTNQ
19702	50070	В	19817	61	969	
19703	50071	A	19818	60	436	RGSREYLHTGCPRSGGCRSQCP
			_			GPPTQSPKVRPPQRGAPRLRGG
						DFPPSRLLDTVSFPSPA*AAFPP
						CPVPHLPAPVQGAPREVVDPSG
						GWATLLESIRRAGGIGKAKLRS
						MKERKLEKKKQKEQ
19704	50072	В	19819	575	1666	``
	<u> </u>				I	<u></u>

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19705	50073	A	19820	515	1110	MFHPTCLTCPAFANRPHVHCRP GAPAFAPSAPGTIPELPTFHTEV SPSHSRQTLQDGVT*TAPPPPPP PPPPPPAPEVLGQLQPPLPTLQP RPLLGQGAREGRTAAAA/PSPS/ GPGSSQGSGRPLRWLGHSARV HPPSWGHRQGLAAQHERKRNV EKKKQKEQEQVRATSQGGHLM SDLFNKLVMRRKGHLWEKDPG AW
19706	50074	Λ	19821	1	191	MQK*ITAWAPAPMKIKIIASPER KYSVWIGGSIWPQLST/FQQMW ISKQEYDESGPSIVHRKCF
19707	50075	С	19822	791	970	
19708	50076	Α	19823	13	1260	INPPPLSRRCQLSHSVLPPLRRR VSLPVAMEEEIAALVIDNGSGM CKAGFAGDDAPRAVFPSIVGRP RHQGVMVG\MGQK\DSYVGDE AQSKRGILTLKYPIEHGIVTNW DDMEK\IWHHTFYNELRVAPEE HPVLLTEAPLNPKANREKMTQ\ IMFETFNTPAMY\VA\IQAVLSL* \ASGRTTGIVMDSG\DGVTHTV\ PILRGLPHCSTPFLRLGPGLARD LTDYLMKILT\ERGYSFTTHGPS GKTFRNIKGEACATSPLDFEQ\E MGT\AASSSSLEKSYELPDGQVI TIGNERFRCPEALFQPSFLGMES CGIHETTFNSIMKCDVDIRKDL YANT\ALSGGTTMDPG\IADRM QK\EI\TALAAPAP*KIR\IIAPPER K\YSVWIGGSILASLSTFQQMWI SKQEYDESGPSI\VHRKCF
19709	50077	A	19824	93	534	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	•	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			:	sequence		
19710	50078	Α	19825	1	1462	WEVSAQPSPAVPRRAPLRGLSA
						TRSAAGTMREIVHIQ\AGQCGN
						QIGAKFWEVISDEH\GIDPTGSY
:						HGDSDLQLERINVYYNEAAGN
						KYVPRAILVDLEPGS\MDSVRS
İ		İ				GPFGQIFRPDNFVFGQSGAGNN
						WAKGHYTEGAELVDSVL\DVV
						RKESESCDCLQG\FQLTHSLGG
						RHGVPGWGTLLHQQGSGEEYP
		-				DRIHETPFSVH/VPSPKVFKTRL
İ						LEPLQRHTSRFHHLVENT\VKP
		1				YSIDNEALY\YICLPTAES*PTPT
						YG\ALNPPGCSATMSGVTT\CLR
						FP/GGQLNADLRKLA\VNMVPF
						PRLHFFMPGFAPLTSRGSQQYR
						ALTVPE\LTQQMFDSKKMIA/AP
ŀ				}		CDPPPTGR\YLTVAAIFRGRMFH
						GRRWDEADGSNVAGTRNRQVI
						SWEWVP\NNVKTA\VC\DIPPPG
						AWKMS\ATFIGNQPRPSRRLFK
						RIYE\QF\TAMFPAQGPSLHWLH
ŀ						GRRAMDGDGSSPEAREQHERP
						WCPEYQQYPGTPRPT
19711	50079	Α	19826	2	180	
19712	50080	A	19827	28	185	
19713	50081	Α	19828	179	307	
19714	50082	Α	19829	438	663	
19715	50083	C	19830	99	335	AVADOL *I DDINGGGGGGGGGG
19716	50084	Α	19831	1	320	AYADQL*LRRHKSSSSSSSSSSSSSS
ŀ						SSSSSSSPLAP*LGSGDLP*EIN
						PPSSCSLLREKDPPTTSGPQ\PTS PRNISPISNPDPTGNRTVQLTWQ
			ļ			,
10717	50085	A	19832	578	753	PLPQPLELWPKAL LCTPTPFPLFWKTVRKYSSNNQ
19717	30083	A	19032	378	733	KGE*NTVLMEVCPRSPRTN*KD
						SQ*PFELLTKMTSQGSDISGDLP
						WEINPLSSCSLLHEKDPPTTSGP
						QTDQPKKHLTNFKSETKETHFI
	:					RGPKTPVLVTDWEGRLPLVFN
1						HSRDASLIIHPRFRGVRPRRDAC
						LGPSPLAASPAFLEEGQVPQPLL
						SMSLTPSLLFWRRGKKPSTPSSP
1						LAASPAFLEEGQVPQPHISVPRS
						LISTPOPLITLHPDPFPAFLEDSK
						EVLKQQPKGRIKHCADGGMSK
					1	EPKNQLKGLPVTKAGNIHGIND
19718	50086	A	19833	95	259	SHRPSQ*VLQLIKVAHPELFVPP
19/10	150000	\ \frac{1}{2}	1,7033			SGFVVSLTSGMKLQTLTTREW
						MRILLIFWNG
19719	50087	Α	19834	1	939	
19720	50088	A	19835	391 .	792	
19/20	120000	14	17033	1371 .	1,72	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19721	50089	Α	19836	1	1317	
19722	50090	В	19837	1	1665	
19723	50091	A	19838	573	902	YSCLLILIRKITQNSVEVLGRRK FLGGGMEREWVMFL/EGCFKR D*GQRGNQFQIRVTAPNFMCV
						CVKRPPNRLCVSNKAVYFTWV QVGFQIFEKPEPIEEICKVCWRP
19724	50092	Α	19839	1802	2229	GGGAEAEEELGPSSAWRGAAC PCPRKAGLVAKAEGPRQASLR GLTPLKRECIIREASLQ*LNTK/V KAAFPVRDRRWSFGFTDKTCLF CLGSEIGEVFLGLVGLRT*GRR WIFLMEQRAGGQGIDLPREVPR SKSGHQISLVSI
19725	50093	A	19840	19	4153	PPDPLPGLPCPPGGPPLPAFGGG WGGARGSWHWSSRACSRRRR LVHAPRAPLLPRAAAEKAKRP AGARQMGLKARRAAGAAGGG GDGGGGGGGGAANPAGGDAAA AGDEERKVGLAPGDVEQVTLA LGAGADKDGTLLLEGGGRDEG QRRTPQGIGLLAKTPLSRPVKR NNAKYRRIQTLIYDALERPRGW ALLYH\ALVFLIVLG\CLILAVL\ TTFKEYETVSGDWLLLLETFAIF IFGAEFALRIWAAGCCCR
19726	50094	A	19841	355	640	ILDYVFSSSHLFQSKLLGRRGR ASHGTKL*AGPG\GGGEVIKGL* GGGAEAEEELGPSSAWQGAAC SFPRKAGLAAKGEGPRQASLH GLTPLKCG
19727	50095	С	19842	55	189	
19728	50096	В	19843	96	401	
19729	50097	С	19844	558	725	
19730	50098	С	19845	448	540	
19731	50099	С	19846	140	334	
19732	50100	A	19847	362	710	ILGGRKLRLEKSLGCAAQGGG AL/YEEWLELRNGCLCCS\VKD NGL*/RAIENLMQKKGKF\DYIL LETTGFADPGAVASMFWVDAE LGSDIYLDGIITIVDSKYGLKHL AEEKPDGLIQ
19733	50101	Α	19848	181	543	
19734	50102	Α	19849	1	594	
19735	50103	В	19850	29	746	
19736	50104	A	19851	45	362	EEQHSMLGSGFKAERLRVNLR LVINRLKLLE/KKDERARIRVEH IIREDYLVEAMEILELYCDLLLA RFGLIQSMKELDSGLAESVSTLI WAAPRLQSEVAELKIVA

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19737	50105	Α	19852	344	951	EEQHSMLGSGFKAEHLRVNLR
	-					LVINRLELL/EGKKKKKTELAQ
1				Ì		KAGKEIADYLAAGKDERARIR
						VEHIIREDYFVEAMEILELYCDL
						LLARFGLIQSMKELDSGLAESV
		ŀ				STLILAASRLQSEVEGFPNIYSS*
		Ì	1			IPSNSPIILNL*MAIKGCKNY/RL
				1		LHQIVVPGPSQKPLQNFLLDLR
·						LPWTTLWPIFGLEGKIPLWERIK
19738	50106	A	19853	1	1077	
19739	50107	Α	19854	1	1140	
19740	50108	Α	19855	1	3426	
19741	50109	Α	19856	1	942	
19742	50110	Α	19857	342	1499	EEQHSMLGSGFKAERLRVNLR
						LVINRLKLLEKKKTELAQKARK
						EIADYLAAGKDERARIRVEHIIR
						EDYLVEAMEILELYCDL/LCLA
						RLG/LYSSY*KELDSGLA\ESVST
1						LIW\AAPRLQSEVA\ELKIVADQ
						LCAKYSKEYGKLCRTNQIGTV
						NDRLMHK\LSVGTPP\QILVERY
						LIEIAKNYNVPYEPDSV\VMAE
						APPGVETDLIDGGFPDDVKKR\
						GPWKKEGSGGFTAPVGGPDGT
						VPMPMPMPMPMPSANTPFSYP
						LPKGPSDFNGLPMGTYQAFPNI
						HPPQIPATSPIRIESVRWTFNCLI
			1			KEYS/IFLAQICLGPGPKARRPF
						AKFP\SRPADNF*QLLFLPELPSV
:				-		PDTLPTASAGASTFRHLKDIDF
10510	70111	<b> </b>	10050	1	0.55	DDLSRRFEGAGKRKT
19743	50111	A	19858	2	255	GCISTSRLGSMPTTTLRLLQKIS
			}			*RSYSSAPASCLISIVKSASD*RA
						RAPTRGSYGGPLLRLSSSSSLAS
10744	50112	<b> </b>	10050	1	276	GRCVTSRASLRPRS
19744	50112	A	19859 19860	3	276 151	GAVPAVRLGGPLHVHLPHAR*P
19745	50113	A	17000	[]	121	SNLSALQFYQDGSLRWWQTVS
						*RAASRSPPTRPVTIKSVCTPILP
						GWLSSLVANCIVKSASD
19746	50114	A	19861	1	386	DCVIATDAEVLGAPVKGKRRG
19/40	30114	^	17001		300	RGPKGDKASDPEAGVARGIDF
						HHVSAVLNFDLPPTPEAYIHRA
						GR**CDGPGICMVGTLRDLGCA
						GQSLRVGDASCTVFPCRTARAN
						NPGIVLTFVLPTEQFHLCKI
19747	50115	В	19862	26	186	NI GIVE IT VET TEQUILECKI
17/4/	120112	۵۱	17002	120	1,00	

				Nucleotide		Amino acid sequence (X=Unknown,
	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		07540,217	sequence	or pepular sequence	section, ( possible nucleotide insertion)
19748	50116	Α	19863	2	1839	ARGQRVRAALLSSAMEDSEAL
						GFEHMGLDPRLLQAVTDLGWS
						RPTLIQEKAIPLALEGKDL\LAR
						ARTGSGKTAAYAIPMLQLLLHR
						K\RQVRW*NRQ*EGLVLVPTKE
						L\ARQAQS\LIQQLATYCCSGMS
						ELANVSAC*KTQSLRELCLMEK
						PD\VVVGTPS\RILSHFAARQPET
						FVTPWSFLVVERSLTFFFPFGFE
						EELKKSSSGKAESLAPDFNKAF
İ						LMSATFNEDVQA\LKELILHNP\
						VTLKLQESQLPGARPVTAVSGG
						L*DLREDKFLLLYALLKLFIDSG
						ASLWLFCSTL*EREFTGLRLFL
						WEQFQ/LSPTCVLNGELPLRSR
						VPHHL/QQFNQGFYGLCHRQL/
						NAEVLGAPVQGQASGPRAQRG
						Q/VPLDPEAGVARGIDFHHCVC\
						VLNF*SFPPTPEA\YIHRAGR**C
						DGPGICMDSTR*QPRA*S*PFVL
						PTEQFH/CRQD*GASQWREQGA
						P/SLLPYKFGMEEIEGFRYR\*GV
						HPQDAMRSVTKQAIREARL\SLI
						KEELLHSEKLK\TYFGRQPLGD
						LQA\LRHGPYLCTPQLVEAPTW
						GHVPDLPGSSCFSVALGAPFTK
	i					KREEACLPSCRKAQESKVPRTP
						LRSFKHKGKKFRPDSQALREVV
19749	50117	Α	19864	3	378	TTTKFAGRRPKVTVLGQP/KAA
						PSVTLFPPSSEELQANKATLVCL
						ISDFYPGAVTVAWKRGSSPVKA
l						GVETTTPSKQSNNKYAASSYLS
						LTPEQWKSHKSYSCQVTHEGST
						VEKTVAPTEYLLRVY
19750	50118	A	19865	1	546	MAWTPLLLLLLSHCTGSLSQPV
						LTQPPSSSASPGESARLTCTLPS
						DINVGSYNIYWYQQKPGSPPRY
						LLYYYSDSDKGQGSGVPSRFSG
						SKDASANTGILLISGLQSEDEAD
						YY/CGCFTVFWSTF*VLVPEHQ
						QSWLPLWTPGVVSLLV/CLVPS
						FVPPPLTKVPSSLSSPVKESAILA
						VDIH

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19751	50119	A	19866	2	838	RVRVSTMAWTPLLLLLLSHCT
						GSLSQPVLTQPPSSSASPGESAR
						LTCTLPSDINVGIYNIYWYQQK
						SGSPPRYLLYYYSDSYKGLGSG
İ						\VPSRFSGSKDASANTGILLISGL
i						QSDDEG\DYYCATWDSSLTGQ
						VFGGGD/TKLTRP*GQPKACPP
		1				GVTLVPNPPS/SRSFQAQQGPQL
						VMSS*SDFLTPGSPLNSCPWK\A
						DSQPPFKAG\VETTHTPPKQSNN
<u> </u>						QVRGPAGYLSLTA\EQWEVPTE
						SYSCQ\VTH*RGAPWEKTVGPY
						RMFHRFLNPSPPHHGMT
19752	50120	Α	19867	1338	1854	NSETREHNHRYVPQRETSGGTD
						GWKSKQPE*L*RRQLVQSWAK
						S*NQPKLPSSPH\*ELPWVTPSA
						RNFRSGPYSPPPPPPPQPLPSSHL
						SSLHFYSPFVSYTSFDLYSSSYH
						SYLMDEDTVEPKELRSSQKVIG
		ĺ				HSFTGFQPTLSANPYDRGDGKII
						ELLPRKAQTFPVVRT
19753	50121	Α	19868	1	500	GSTYSELLSGTACVVTALHLAQ
						QRRHHLLGGLASRHDAPEQLA
						VGAPVQPRPLPAPAGAQALVG
						HAHVVVEGGP/PGSGGPGEGAI
						ARGPGAAAAAGAGPGAPRRRR
						LQASAGPCSRSSEPGRRSAPRA
						APPAPSAGPHVAFPSRRPAAGS
		<u> </u>				PPRSRPPPPPGIPTGGR
19754	50122	Α	19869	20	507	TRQVYSGGVRTPPCQSHPHRPR
						SSCRPPPPPRRPRSSRRRRGLSP
						RRAPAHWSSRSTRPAWWAPSR
						RHAPRPPWSPGPASG*CSVYFPI
						LPLALFPLPVHHGGVHVCGREG
				1		VGLVQEGDSHSGEWS*RSAWG
						STSLRAAPRSGGRPPAGGGWR YTRPPSRRC

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19755	50123	A	19870	236	1191	GSGRIPGRAGLEGPGPAFQKSR
' ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' '	100125					LSPASGSAGPRGASPSAVPPGR/
						RPGVPAFPASNPPPVAAVPGPW
						VTCWGKGLQGEGKAPRAPHA
						GTPQR\SPGSARPTGLSPPA*PSR
						*RARRPSPETP\PGHGRAGPPPP
						DGAVQRQGPGCRAPSASRPPGS
						ARARRRWPAARPPGCPARPG
						PRGGRPRRRPSPPAAGRARRRA
						PRTPAARSR*VCPRGRRSAAPP
						RGHGTAPCAAAAPPPPRGLGRP
						ADAPEO/PRCRCARPAPPLPAPA
		1				GAQALVGHAHLLLKAALGHDF
		ł				TFLWATIHEVIPKMRWGQSQC
		ļ				QRGGPSSHTPDLSA
19756	50124	Α	19871	24	286	LRGQKVLLLGDRARLSQKKKE
		l				RKREREREGHGRNGR/RRNG
		l				QGRGPEGQR*NERTGQERTGH
						GREGRKEGRKEGRNERE
19757	50125	Α	19872	43	373	PRFTDENLEMCTVIQSVQCYHR
						SWH/HSSPPEPAPHHQSPPAHSP
						VLCWFRRLGGGEDWAPSSGGR
						AVWRSHCSCFCCSSSCCSCSQ
						SGKRTAAAPWPTTSPAPSRSCC
		l				A
19758	50126	Α	19873	3387	4083	AGRSALPVDAVQVQAEGLWPP
						АVLLНННННРНННҮННҮНН
						ННННННР*НННРННННННRN
		l				ннининининининин
		l				YHSH\YННН/ННННDНННҮНН
			1			FHDHYHHHHH/HYSHH/HHPH/
1						НННН*КНОНУ/АНННННО\НН
						СННСННРНКНСНҮНҮНННСТ
		l			1	VAFGALEDSAVARTGLPLPEGV
						LCGAPTAAASAAVPPPAVPAPN
		l				QGRGPQLHPGQQLRPLLHAHA
						ALQWPTTHL
19759	50127	A	19874	3	411	
19760	50128	Α	19875	1	1095	
19761	50129	Α	19876	1	471	
19762	50130	Α	19877	1	554	FRGRLASPFRPVPMEALGKLKQ
1		1				FDAYPKTLEDFRVKTCGGATV
						TIVSGLLMLLLFLSELQYYLTTE
1						/VKGRGLMRGRGLAF*E*DL*G
		•				WEWGRRGLGLSLTEVALPQVH
						PELYVDKSRGDKLKINIDVLFP
						HMPCAYLSIDAMDVAGEQQLD
		1				VEHNLFKQRLDKDGIPVSSEAE
						RHVLSFVVSAD
19763	50131	В	19878	1116	1274	
19764	50132	Α	19879	28	389	

SEQ ID		1	SEQ ID NO:	I		Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19765	50133	Α	19880	75	448	
19766	50134	Α	19881	276	830	QEDSCPLHPLSGIPFRRFQPTPE
						EAGAVVPGEGPTHDQGPSSALT
		1				RR*ACQGPLHLARTTAEPGHAH
						PWH*RWQRYMGKLGRDLEKV
						CPVWSSVACSSPGLSWAGKTE
						AKTSLEGGNSASASASSWAWG
		ŀ				RSSLAPVRDILGSREFSGLLEGT
						FVLSVTLSADFGIGTWVDLGKV
		ļ				TGSSSKEGRL
19767	50135	A	19882	223	1269	APVTKRKEVFAKDSKGSALDA
						GRDPKRPALPETLCESGWASNT
						APTAPPQPGWCLCGKDFKSSCQ
						TPGREKERRLATMHGSCSFLML
						LLPLLLLVATTGPVGALTDEE
			:			KRLMVELHNLYRAQVSPPASD
						MLHMRWDEELAAFAKAYARQ
						CVWGHNKERGRRGENLFAITD
						EGMDVPLAMEEWHHEA*/DTT
		ł				TSAP/HSCSPGQMCGRYTQVVW
						AKTERIGCGSH\SCEKLQGVEET
		1				NIELLVCNYEPPGNVKGKRPYQ
						EGTPGSQCPSGYHCKNSLCEPI
						GSPEDAQDLPYLVTEAPSFRAT
		ł				EASDSMKMGAEGPDKPSVVSG
		]				LNSGPGHVWGPLLGLLLLPPLV
		1				LAGIF
19768	50136	Α	19883	2	281	FVPALGSSAVMLSRSRCVSRAF
						SRSLSAFQ/KGVSLCQGPGYPNS
						RKVVINNSVFSVRFFRTTAVCK
						DDLVTVKTPAFAESVTEGDVR
			:			WEKGKI

SEQ ID			SEQ ID NO:	1		Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	Sequence		07/340,217	sequence	or peptide sequence	detection, (~possible nucleotide insertion)
19769	50137	Α	19884	1	1320	MPAVTILKPNLKQIILGLKAAE
						VSSLPPSIVLDAKEITTQITRQIL
						KGNCPLGRHSLPGVSLCQGPGY
		İ				PNSRKVVINNSVFSVRFFRTTTV
						CKYDLVTVKTPAFAEPVTEGD
						VRWEKAVGDTVAEDEVVCEIE
		l				TDKTLVQVPSPANGMIEALFVP
						DGGKVEGGTPLFTLRKTGAAP
						AKAKPAEAPAAAAPKAEPIAA
	,					AVPPRAAPIPT/QMPPVPSPSQPP
						SSKPVSAEAQNTCA\MLTIFNEI
					-	DVSNIQKMRARHKEAFLKKHN
i		ŀ				LKLGFMSASVKASAFALQEQP
					:	VVNAVIDDITKEVVYRDYIDIS
						VAVATPQGLVVPVIRNVEAMN
						YADIEQTITELGEKARKNEFAIE
1				i		DMDGGTFTISNGGVFGSLF/GTP
						IINPP\SAILGMHGIFDKPVAIGG
						KVEVQPMMYVALTYDHRLIDG
						REAVTFLRKIKAAVEDPRVLLF
19770	50138	A	19885	1	1559	MESEALVVCINHCMAVNYGMI
		ŀ				YQRGSSNMNILSLSLKHPEKTGI
						SPTANHGISDLQEGEAACGSLR
						TPLVGRRVLRPAPLPVVVRPYI
						RCPPALGSSAVMLSRSRCVSRA
	]					FSRSLSAFQKGNCPLGRRSLPED
						DLVTVKTPAFAESVTEGDVRW
				İ		EKAVGDTVAEDEVVCEIETDKT
						SVQVPSPANGVIEALLVPDGGK
į						VEGGTPLFTLRKTGAAPAKAKP
						AEAPAAAAPKAEPTAAAVPPPA
			:			APIPTQMPPVPSPSQPPSGKPVS
						AVKPTVAPPLAEPGAGKGLRSE
						HREKMNR\MRQRIAQRLKEAQ
						NTCAMLTTFNEIDMSNIQEMRA
						RHKEAFL\RKLKLKLRLMSAIV
						EGSA\FALQE\KPLLMEGIDDTT
			1			KEVVYRDYIDISVAVAHPRGLV
						VPVIRNVEAMNFADIERTITELG
						EKARKNELAIEDMDGGTFTISN
						GGVFGSLFGTPIINPPQSAILGM
						HGIFDRPVAIGGKVEVRPMMY
				1		VALTYDHRLIDGREAVTFLRKI
					_	KAAVEDPRVLLLDL
19771	50139	В	19886	239	3045	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	İ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19772	50140	Α	19887	3	809	RGRFLPPCHPAALPPCRPAALLP
						VVAARGAPSPPPPHVLPCVGSG
		l			İ	LGYNMLGMIKNSLFGSVETWP
	1					WKVLSKGDKEEVAYEERACEG
						G\KFATVEVTDKPV\DEALTGK
		İ				QLPKV\AKYAGGTNDKG\IGMG
						MTVPISFA\VFPN*RWLSCRKKL
						KVWFPDFQTNFQSDPPAPS\DKS
						VKI\EEREGIHCLIPCQFGGYAK\
						EADYV\AQATR\LRAALGG\PAT
						YRGDIYFCTGYDPPMK\PYGRR
						NEI\WLFEDMSDPTEPRTYWEV
						CLCVSFLRG
19773	50141	С	19888	164	208	
19774	50142	Α	19889	1542	1929	NLVPSLGSGDLPWEINP/LSSCS
						LLHEKDPPTTSGPQTDQPKKHL
						TNFKSAARPT\SWGRGKYPSTPS
						PSPLAASPAFLGGKNPPIAYFHT
						PTSYLCAPIPYFCAPTPSLLFWR
					ARTTTPTPSLCLKSFL	
19775	50143	Α	19890	1	208	MPEPQRPGVPPEPPPGACYAC
į	1					RKSGHWA\RNARSPGFLLSRVP
						SPPGPSRTPSFG*LSQWKPTCTQ
						VK
19776	50144	Α	19891	1	1662	
19777	50145	В	19892	1	915	
19778	50146	Α	19893	183	569	
19779	50147	C	19894	71	344	
19780	50148	A	19895	147	446	
19781	50149	A	19896	167	373	
19782	50150	A	19897	142	363	
19783	50151	C	19898	52	174	
19784	50152	C	19899	228	260	
19785	50153	A	19900	1	753	
19786	50154	A_	19901	1	1077	LGSGDFPWEINPLSSCSL/LREK
19787	50155	Α	19902	500	912	
						DPRTTSGPQTDQPSKHLTNLKS
						ASTPPPYNPFITSPPHTRSGLQFR
						STSSPPAPAQQFTLKKVAEAKG
						IVKVNAPFSLSDLSQISVRLGSFI
10700	60156	<u> </u>	10002	1,	061	KYEKSSPVHGSFGSNPETLYSP
19788	50156	A	19903	174	861	
19789	50157	В	19904	174	222	
19790	50158	Α	19905	1	756	

NO: of p	peptide l quence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)  NLVPSLGSGDLPWEINP/LSSCS  LLHEKDPPTTSGPQTDQPKKHL  TNFKSAARPT\SWGRGKYPSTPS PSPLAASPAFLGQRQVPLSWGR  GRCPSTPSPSPSAASPAFLGQGQ
19791 50			,	sequence		NLVPSLGSGDLPWEINP/LSSCS LLHEKDPPTTSGPQTDQPKKHL TNFKSAARPT\SWGRGKYPSTPS PSPLAASPAFLGQRQVPLSWGR
	0159	A	19906		1852	LLHEKDPPTTSGPQTDQPKKHL TNFKSAARPT\SWGRGKYPSTPS PSPLAASPAFLGQRQVPLSWGR
	0159	A	19906	1035	1852	LLHEKDPPTTSGPQTDQPKKHL TNFKSAARPT\SWGRGKYPSTPS PSPLAASPAFLGQRQVPLSWGR
			19900	1033	1002	LLHEKDPPTTSGPQTDQPKKHL TNFKSAARPT\SWGRGKYPSTPS PSPLAASPAFLGQRQVPLSWGR
19792 50						TNFKSAARPT\SWGRGKYPSTPS PSPLAASPAFLGQRQVPLSWGR
19792 50						PSPLAASPAFLGQRQVPLSWGR
19792 50						
19792 50						OKCI STI SI SI SAASI AI LOOKKI
19792 50						RQVPLSWGRGRCPSTPSPSA
19792 50						ASPAFLGQGQRQVPLSWGRGR
19792 50						CPSTPSPSPSAASPAFLGQGQRQ
19792 50						VPLSWGRGRYPSTPSPSPSAASP
19792 50					į	AFLGQGQRQVPLSWGRGRCPS
19792 50				ŀ		TPSPSPSAASPAFLGQRQVPLSW
19792 50						GRYPSTPSPSPSAASPTFLGQDL
19792 50	1					PLSWG
19792   30	0160	_	19907	1644	1981	LGSGDLPWEINPSSSCSLLREKD
	0100 1	A	19907	1044	1981	PPTTSGPQTDQPKKHLTNFKSA
						*KRTNTASYK/WHSQSRVPAPR
						DGLVAAPTESEVAFLPVPPHQG
						PPLGNLGPSTVTTSFRCWLLDP
	1					
10702 50	0161		10000	1011	1154	RS
		A A	19908 19909	1043	1168	
		A	19909	1328	1470	
		B	19911	406	1191	
		A	19912	2848	3140	SFYHLSSSHLVRLTVSFRD*PSP
		^	17712	2040	3140	TCPAIYS*KGGWSQRHSQGACY
						TCRKSGHWA\RNARSPGFLLSH
	1					VPSVWDPTENRTVQLTWQPIPE
						PLELWPKAL
19798 50	0166	В	19913	1	941	
		_ A	19914	656	1983	NLVPSLGSGDLPWEINP/LSSCS
	j				†	LLHEKDPPTTSGPQTDQPKKHL
						TNFKSEAKEICFIREPKTPAPVT
		l				DWEGSLPLVFNHCRDASLIIHS
						RFKGVRPRRDACLGPSPLAASPI
						FLGKGQRLKTDTARLPRKPPRP
	•					SRMPSFGVKHVNVTIDCLPEGA
						ATRGTARTSKPTTKSQKTLPST
						SPGHWTQSTPWASALRSSPWTE
						TAAPSETEETLNTGRPPELPARA
						TATWFSASHTLPALATRRVART
						QWLTADRQTWASISSVPWAQT
						ISEKKPGGSLWETRSSPPTTAGT
						EEAMNTTSLLAPAAEIMATPGS
						PSQASPTSGAFTHGTQTPSPTKA
					<u> </u>	TAPRYPOTGNSILTFGRPFLPFA
						GDLFAEWPFTAGEEPVLVPRPH
						QVSKTRSPPSFHLTLKTVTVAL
						LGNGEECYVYLSVSELREIRTR
						LYSPDAGLNEEKMLTKLLLI
19800 50	0168	A	19915	117	432	
L		C	19916	122	289	-

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence ( X=Unknown,
NO:	of peptide sequence		in USSN 09/540,217	location of first codon for peptide		*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,217	sequence	or peptide sequence	deletion, =possible nucleotide insertion)
10000	50170	<u> </u>	19917	1	516	
19802	<del></del>	A	19917	212	3302	
19803 19804	50171	B C	19918	181	444	
	50172	A	19919	151	485	CDD CHVOLLI SCRAL ADDVDDI
19805	50173	A	19920	131	483	SPRGHYQLLLSGRALADRYRRI YTAALNDRDQGGGSAGHPAS/
					1	RSENLENTVIIPDIKLHSNPSAFN
						IYCNVRHCVLEWQKKEISLAAA
						SKNSVQSGESDSDEEEESKEPPI
19806	50174	A	19921	1	3465	3KN3 VQSUESDSDEEEESKEFFI
19807	50175	A	19922	1	176	
19808	50176	A	19923	114	14211	AAPVAAPGALFMPVPDGSVAA
1,7000	30170	``	17723	114		AGLGLGLPAADSPGHYQLLLSG
						RALADRYRRIYTAALNDRDQG
						GGSAGHPASRNKKILNKKKLK
	:					RKQKSKSKVKTRSKSENLENTV
						IIPDIKLHSNPSAFNIYCNVRHC
						VLEWQKKEISLAAASKNSVQS
						GESDSDEEEESKEPPIKLPKIIEV
						GLCEVFELIKETRFSHPSLCLRS
						LQALLNVLQGQQPEVLQSEPPE
					:	VLESLFQLLLEITVRSTGMNDST
						GQSLTALSCACL
19809	50177	A	19924	375	1191	SQGIPELHYTPGPRGRWCGWW
19609	30177		17724		1171	PIVEPRMGASQGCCVLGHLAQ
						QGGGASQAVRGANGVHAGCR
						GDPHPRSDHYPGGRPWLPGAS
						QDPAVAPGGHGHPSYLARLPC
				1		AAEPGTP/PAEPPRPSSLPGAAT
						APAGT*\PALADAPPRRVGGAG
						QPGRGHV*PKLVPGWQSQDSI*
						PSL/PSSPARHTLAASATP*APGS
<u> </u>						GARPAW*C*WAPALAPVIPVD
						AHSPP\GGAGHGPGH*GPRGSV
						LGLRLPAGCP\GPRQPHRPGQSG
						PQPSSPPGGALRSASR
19810	50178	A	19925	3	543	SDDQPEKPHFDSRSVIFELDSCN
						GSGKVCLVYKSGKPALAEDTEI
1			}			WFLDRALYWHFLTDTFTAYYR
						LLITHLGLPQWQYAFT\SYGISP
1						QAKGITKATCEWGEHIIPLGSM
		1				AINSISKLTQLTQSSMYSLPNAP
						TLADLEDDTHEGRKGTEGSQL
		1				KCEHQLPPPMSHVGGGQRESL
						ARKEK
19811	50179	A	19926	1	392	FLDRALYWHFLTDTFTAYY\RL
1,011	1	` `		1		LITHLGLPQWQYAFTSYGISPQ
						AKITYNIKMNRRKCVHWKLAV
		1				EPLQWELRSCVSAGMGTGPKQ
						KVSSMKPRSCSVCSRRQSNSMG
				[		TTVSQSFLSYGKNVLFFPVRCC
L	<u> </u>		<u></u>		<u> </u>	1.10001D01GKH4D1114KCC

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19812	50180	Α	19927	1	1081	RAMEEEASSPGLGCSKPHLEKL
						TLGITRILESSPGVTEVTHEKPP
		l				AERHMISSWEQKNNCVMPEDV
			İ			KNFYLMTNGFHMTWSVKLDE
						HIIPLGEALAINKHLKTDFSFTQ
						SSMLFTFLIAPTLGQTWEDDTH
		1				GSQWMIQPEKPHFGLFAVLIF*
		l			•	GWVS\CNGSG\KVCLV\YKSG\K
					į	PALAEGHWRFLFPGTELLYWH
						FLTDTFTAYYR\LLIT\HLGLPQ\
						WAILPFTSYGISPQAKQW\FSM
		1				Y*TYSPYNTNLV/TQKETDSFCG
						IKLDPSKVF*GGKNKIVIPKKKG
		1				PVQPAGG\QKGPFRDPPVPSISSI
		1				FKSSSGLWGNPTREVSTPPSNSL
				İ		PAPEWWFPCTDGPRGDLQFCV
						WTVGLFLVE
19813	50181	Α	19928	1	2835	MDALMLALGALMLALGLGDP
						KSAQVSDPDGAEPGPGRATGT
						DPLVQIHWKLGISEALSIPGGRL
						YRDLWSLTPVCLHIPGIAHHGP
				ļ		FTLGRMDVVEVAGSWWAQER
						EDIIMKYEKGHRAGLPEDKGPK
		i				SFGSYNNNVDHLGIVHETELPP
						LTAREVKQIRREISRKSKWVKM
		İ				LGEWDTYKNSRKLIDRAYQGIP
						MNIRGPMWSVLLNIEEIKLKNP
						GRYQVRSARAQPTGQAVSGAQ
						VSSWRERQDHPGELGVKI
19814	50182	Α	19929	3	142	TISSIIRKÄRKP\QDLLNNYTPRS
						SPLRSSGRTRMYKGKNVRPGNS
		<u>L</u>				S
19815	50183	Α	19930	904	1295	WALMNGSESRFFVSSSQGRSEL
		1				VPNS/NYAVDY*PLTAHAVSSG
						GTSPVLARLLREKLESL\LPLHL
				}		GQVAKYAGQLRGRVKQQFAT
						MGERRRFWEKLFVNDR/AGAV
						AGKQRSESHY*NDRTVNQRTA
						RPSR
19816	50184	В	19932	1	1050	
19817	50185	Α	19933	303	384	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	E .	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19818	50186	A	19934	375	1161	EALESAPEG/RR*GSGTKTLSPS
						P/PAGSWVCRIPMGRGCRHGAE
						R/P/PGSQESSPSNGHGKLAGPSP
						YLGRFKSSS*T*QSPQTPAQGVS
						AS*GLGAAPVLRLQGPVPP\SHC
						GKSRPDPC*PSPGSPDPPGERES
						QQESQ*WPPVGELCTDPTGNPE
						RGGRHLQ\SHAPSGGGLSGFPS
		1				DSFPCSCYYGSTPVARKKRFAK
		ŀ				CRGYIYDFVYELLSLKPPSQGV
		ŀ	İ			AEMPPPAVGQQG\QAEPRGIPQ
						SPAGGRVFFQDFYSLCLENTS
19819	50187	В	19935	62	346	
19820	50188	В	19936	1	1629	
19821	50189	Α	19937	1	1218	
19822	50190	C	19938	718	1053	
19823	50191	Α	19939	54	212	
19824	50192	Λ	19940	122	451	
19825	50193	Α	19941	329	398	FFSYFRPSGCILPL*ELPEARCP
19826	50194	Α	19942	92	346	VGSSPSRL*SLRGQVLNKN*S/I
						QLGCLHIVGNPI*NS*DVPWAG
						WSED*GRRWISSRSSSKDRGPV
	1					S*RSPPVPGLQHQMSHASM
19827	50195	Α	19943	284	374	LKNHKRSEYALPHLN**PSTTK
		İ				EV*M/DRS/SALSDDITL*KSFSW
						LILAQKSPPLSTLQPPLLPAREQ
						TLFDCNSPLPTQIL*NGPTLISLR
			_			*LSQDEPGKGLSQGNVIT
19828	50196	Α	19944	1	1674	MAPPLSPFADSLFGLSPPAPRDK
						GDTFYPWTQNSGARHGLGRQP
						SLGVSSLQGRLSDHSPTFQRCQ
ĺ						TTQGRLPWSFTLSGKSCFSEEG
				i e		AIPTPSVAESFQSSFSTDPSDLSP
						PPQPAHRQAELSPNSSSASTPPP
	1					YNPPTTSPPHTRSGLQFSSATSS
						SPPAQQFPLREVAGAEGIVNAH
				•		VPFSLSDLSQISQHLGSFSSDPT
						KYIQEFRYLTLSYNLTWSDLNV
						ILTSTLSPDERERVFPVSQSHAD
						NRRLHEP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19829	50197	A	19945	1746	3501	ESDPSNSTTSMDRTLL/MSSIVP
		1				RLPSACRILPTGFTIPE*SCVHRT
		1				ASLIPPLPPGSRKYSPPLPLNSLI
						FLKN/SAYTAVPALQTDWATSP
:						ISLHLRTSFNSPHLYPPEELIYFL
						DRSSKTSPDISHQQAAALLRTY
						LKNLSPYINSTPPIFGPLTTQTTI
						PVAAPLCISWQRPTGIPLGNLSP
			-			SRCSFTLHLRSPTTNINETIGAF
						QLHITDKPSINTDKLKNISSNYC
						LGRHLPCISLHPWLSSPCSSDSP
						PRPSSCLLIPSPENNSERLLVDT
						RRFLIHHENRTFPSTQLPHQSPL
						QPLTAAALAGSLGVWVQDTPF
-						STPSHLFTLHLQFCLAQGLFFLC
						GSSTYMCLPANWTGTCTLVFL
		l				TPKIQFANGTEELPVPLMTPTQ
		l				QKRVIPLIPLMVGLGLSASTVA
-		1				LGTGIAGISTSVMTFRSLSNDFS
		ł				ASITDISQTLSVLQAQVDSLAA
						VVLQNRRGLDLLTAEKGGLCIF
		İ				LNEECCFYLNQSGLVYDNIKKL
						KDRAQKLANQASNYAEPPWAL
						SNWMSWVLPIVSPLIPIFLLLLF
						GPCIFRLVSQFIQNRIQAITNHSI
		l				RQMFLLTSPQYHPLPQDLPSA
19830	50198	Α	19946	1392	1676	GFCILHIVLEPWFSAPSAPLSTSL
	-					YWLF*LYILLNFFQSFQLLCLW
						FECPPVAQSNLIV*SLLLSARQS
						HSPSSFFPLLVRNCVPLEEERCS
19831	50199	В	19947	1	1500	
19832	50200	Α	19948	107	691	TTMLLIQSLFGGLFTWTRMKFG
						AVARIGGHPLGDQSP\SSCSLLP
	İ	1				EKDPPTTSGPQIDQPKKHLTDF
						KSGTPL*LYTHVSRVSDHAGTP
	:					ALVLHP\YGKSRFSGEGATQRQ
		ļ				YPIPQQALKGLMPAITRLLQHG
		1				LLKPINSPYHSPILPVLKPDKPY
		1				KLVQDL\PLSTKLFCLSTPWCQT
				<u> </u>		HILFYPQYLPLQPIILF
19833	50201	В	19949	1	2628	
19834	50202	В	19950	1	585	
19835	50203	В	19951	178	462	
19836	50204	C	19952	1	825	

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
10027	Isaacs		10052	Z 4 4	1705	PROVIDENTIAL NOVEMBER
19837	50205	A	19953	644	1795	RPSKSQTMLNSLSTALIISKIYFL
						PHT*HIYFLLPGSFCYTHSLLSLP
						QLPLFL/CLDLNAASHIIPDTTPD
						PHDCISLIHLTFTFPQISFFPVSHP
						DHTWFIDGSSTRPN\TTHQQRQ
						AML*YLPHLSLRLPLCLPPLPLS
						KPN*LP*LKPINSPYNSPILPVQK
						PDKSYRLVQDLRLINQIVLPIHP
		l				VVPNPYTLLSSIPSSTTHYSMLD
					:	LKDAFFTIPLHPSSQPLFAFTWT
		1				DPDTHQSQQLTWAVLPQGFRD
		1				SPHYFSQALSHDLLSFHPSASHL
						IQYIDDLLLCSPSFESSQQDTLL
						LLQCLFSKGYRVSPSKAQISSPS
						VTYLGIILHKNTHALPADRWPS
						CLRAAAAAALILVEALKITNYA
						QLTLYSSHNFQNLFSSSHLTHIL
						SAPRLLLLYSLFVESPTITIVPGL
						DLNAASHIIPDTTPDPHDCISLIH
		]				LTFTFPQISFFPVSHPDHTWFID
						GSSTRPNHHSPAKAGYAIVSST
E		}				SIIEATALPPSTTSQQAELVTLT
		ļ				QALTLAKGLRINIYTDSKYAFHI
						LQHHAVIWAERGFLTTQGSSIIN
						ASLIKTLLKTTLLPKEAGVIHCK
						GHQKASDPIALGNAYADKVAR
						QAASSPTSVPHSQFFSFASVTPT
						YSPTETSTYQSLPTQGKWFLDQ
						GKYLLPASQAHSILSSFHNLFH
						VGYKPLDRLLEPLISFPSWKSIL
						KEITSQCSICYSTTPQGLFRPPPF
						PTHQARGFAPAQDWQIDFTHM
19838	50206	A	19954	52	641	RPSKSQTMLNSLSTALIISKIYFL
				:		PHT*HIYFLLPGSFCYTHSLLSLP
						QLPLFL/CLDLNAASHIIPDTTPD
						PHDCISLIHLTFTFPQISFFPVSHP
						DHTWFIDGSSTRPN\TTHQQRQ
						AML*YLPHLSLRLPLCLPPLPLS
						KPN*LP*LKPSLLQKDYASISILI
						LNMPFIFCSTMQSYGLKEVSSL
						HKGPPSLMPL
19839	50207	A	19955	3	380	TTPDPHDCISLIHLTFTPFPHISFF
17037	30207	<b> </b> ^\	19933	3	200	RVPHPYHIWFIDGSSTRPN\ATH
						QQRQAML*YLPHLSLRLLLCPP
						PLPLSKPN*LP*LKPSLLQKDYA
						SIFILTLNMPFISCTTMQEVSSLH
L	<u> </u>	<u> </u>				KGPLSLIPL

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19840	50208	A	19956	154	485	TFIRTHFLTQPQPCPRHQPSR*L SSSPPAG*TG/IFARLLIFSCLLQI LNHM/QHTLAGQSVLVKNLTP QTLQPRWTGPYLVIYSTPT/VRP PAGSSPLGSPFQSKAVSISQPA
19841	50209	A	19957	158	416	ISGTCSNLLGEEVLEERLAAAV QAFGSSRVLEMWLGFVSQWRQ VIVTQKCIAVWLLPLYYCTP*R QG*LIPVVGFEGQILIFEVFS
19842	50210	С	19958	1	1755	
19843	50211	Α	19959	2	394	
19844	50212	A	19960	710	1093	KEPTQHSFSYTHSLLSLPQLPLF L/CPDFNPASHIIPDTTPDPHDCI SLIHLTFTPFPHISFFPVSHPDHT WFTDGSSTRPN\ATHQQRQDM Q*YLPHLLLTLPLCPPPLPLSKP NQLP*LRPSLLQKG
19845	50213	С	19961	1	1287	
19846	50214	A	19962		1293	MEEPLGFKQTLVVAQQNPLWT SGGGGHSPSGSGHRGVCITTLP VPEGSSQKETPHVSEKLKEKNE SLPGNPENSSRSSERPPRRHSKA RDSDEENDPDDEDAVVNAVGC LGKNSGFLAPELQKYQKQIKDG TAGTPAISTTTTVEIRKSSVMTT GDHLLSGKKAQRPAPGHSSCPS TETEEEKGKRLLYWVVWQVA VNSASQLEAHNSENPVSLSCRI RERPGLWHGYRPSRSPLPHSLS GSYSLWALRTDTVCD
19847	50215	С	19963	1	846	
19848	50216	A	19964	564	1172	FPKSIFFLPPDAYTFLFPWLLQL YSLFVESPTITHCSWPRLQSGLP HYSGYHT*PS*LYLSDPPDIHPIS AYFLLSCSSP*SRLIY*WQFHQA *SPHTSKGRLCYSTSH*PAS*NL SFPFHRGNVSSRK*LLSVPSAIL LLLRDYSGPLPSLHIKLRDLPPP RTGKSA\YSTCPESGN*NTSWSR *TLICQS
19850	50218	A	19966	1382	1778	PIGTNTECVVLFQPMETGHSSR VNMSGYK*PC/PPLFGVLSWQN CWRVYPFCLLFGGPLSVMDPE GSPSGC/SNSLGLGADSVTLSTG GVFQPNVRGLNCNGKIVMGRC PLEFVPIPATSPISRKKPERSTAF

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19851	50219	l IA	19967	239	737	RPSFKTPENSPTLVPTWTTLFYA
			.,,,,,,,			LFFSYPHLPTSLIRPKYFNQIICF
						PDYSWSTATSHCCPSSQSK/CLL
						CV/SLTSPQYQPLPQDLPSA*SLP
	ĺ				L*VPTPPLIPLEAALRNIAHSLSI	
	ŀ	į				PPPKNFRRSNTSTLFCFICLINIR
		j				RQECQASEPRPGHRIPCDLHVYI
						QMA
19852	50220	В	19968	1	1321	
19853	50221	Α	19969	1	1069	
19854	50222	В	19970	758	6451	
19855	50223	Α	19971	53	319	CSGSMPSSSSVSRPTSPFGSPLLS
					SPATSSSPKTPPTSATSGSSCPTT	
						GL*MWTWSGFTGPVKAATWR
	ŀ					WTSATYPRWSWRPRAPLCPP
19856	50224	Α	19972	1427	2103	RPVHGHLHAVPVWRDSVRGLA
1						PSSSPRVPGRDLIKQTPRLRARV
					VLSPFKTADAAHSHPGLTQPGR	
		l				GRMKADISAATPGPFMCRVTPL
						ALGVSTEPSPARLVLNFVLFCF
						VLRRSLALSPRLEVLWLDLG\SL
						QPPPP\GFKQFSCLSLPSSW\DYR
		l				CMPPWAG\QFLEF*VEDGV*H\I
		ŀ				GQADLELLTSGDPPASASQSAG
		l				ITGVSN\RAQPQVSECGHGVAL
			:			RRQ
19857	50225	A	19973	3	320	VRMVIIKKSENNRCWRKCGEIG
į						MLLNCW*ECKLVQPVWKT/VD
		İ				PAIPLLGIYPKDYISFYGKDTCT
				-		HMFIVALFTIAKTGNQPKCPSMI
						DWIKKMWHIYTMEYYAAI
19858	50226	Α	19974	2	79	
19859	50227	Α	19975	308	630	GREKAREQVDLTLFTIAKTWN
						QPKCPSVIDWIKKIW/HIYTMGY
					]	YAAIKKDEFVSFAGT\WMKL*T
						VSVEADPHIEGRFFRFKNYASK
						TPSLKLQRQSRPFQETSHPTGIE
						KRSPPNLQKATQPHIMSAKEYA
						FSRKMYGKTGRKKGEVKVRER
i						KGKGAGRSDTIYNSKDLEPTQ
						MPISDRLDKENMDIYTMGYYA
						AIKKDEFVSFAGTMDEAVNQG
				1		YSSRARTRHWSMKKILLQLAR
<u></u>		<u> </u>		1		CLPKWPLSFVLESQGLGGVGS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19860	50228	Ā	19976	2	508	KTTMRYYLTPVRMAIIKKSENN
						RCWRGCGETRMLLHCWFECKL
		ŀ				VQLLWKT\W*FLKDVELKIPFD
						PAIPLLGVYPKDYKSCYYKDTC
						TRMFIAALFTIAKTWNQPKCPS
						VIDWIKKMWHIYTMEYYAAIK
						KDEFVSFAGTWMKLETIILSKL
						SQGQKTKHRMFSLISGS
19861	50229	Α	19977	1926	2448	DFIAVITTRKQLKTKYNAHHSK
						QAITKHFSPKISDLILCLKKKIM
						NRHFSKEDIYAAKKHMKKCSP
	<u> </u>	1				SLAIREMQIKTTMRYHLTPVRM
						AIIKKSGNNRCRRGCGEIGTLLH
		1				CWWDCKLVQPLCKSMWRFLR
						DL\DPAIPLLGIYPKDYKSCCYK
						DTCTRMFIAALFTIAKTWNQP
19862	50230	В	19978	1	1899	
19863	50231	Α	19979	1	2142	MIILIDAEKAFDKIQQPFMLKTL
						NKLGIDGTYLKITRAIYDKPTA
						NIILNGQKLEAFPLKTGTRQGCP
						LSPLLFNIVLEVLAQAIRQEKEI
						KGIQLGKEEVKLSLFADDMILY
						LENPIVSAQKLLKLISNVSKVSG
		1				YKINVQKSQAFLYTNNRQTESQ
		1				IMSEFPFTIATKRIKYLGIQLTRD
ļ		1				VKDLFKKYKPLLNKIKEDTNK
						WKNIACSWIGRINIMKMAFPR
						WELNNENTWTQEGEHHTLGPV
:	<u> </u>	-				VGWGKRGGIALVDIPNVNDKL
						MVLEVLARAIRQKKEIKGIQLG
i		1				KEEVKLSLFADDMIVYLENSIV
l						SAQNLKLISNFSKVSGYKINVQ
ı						KSQAFLYTNNRQTESQIMSEFP
ı		1				FTIATKRIKYLGIQLTRDVKDLF
ı						KENYKPLLKEIREDTNKWKNIP
ı						CSRIGRINIMKMAILPKVIYRFN
1						DIPIKLPMTFFTELEKTTLKFIW
1						NQKRACIAKTILSKKNIAGGITL
1						PDFKLYYKATVTKTAWYWYQ
İ						NRDIDQWNRTEASEVTSHIYNH
İ						LIFYKPDKNKKWGNDSLFNKW
İ				1		CWENWLAICRKLKLDPFLTPYT
ı				1		KIHSRWIKDLNVRPKTIKTLEEN
ı	<u> </u>					LGNTIQDIGMGKDFMTKTPKA
1	}					MATKAKVDKWDVIKLKSFCTA
				1		KETTIRVSRQPTEWEKIFAIYPS
1						DKGLISRIYKELKQIYRKK\TNN
	1					PIKKWAKNMNRHFSKEDIYAA
		1	1	1	i .	I THE HAMMAN THE CINDLE AND
	ŀ					NRQMKKCSSSLVIREMQIKTTM

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19865	50233	A	19981	209	3816	QGRPTFRFRKYREHHKDTPREE
						QLQDT*SSDSPKLK*RKKC*GQ
		ŀ				PERKVKLPTKGSPSD*KRISRQ/
						KTLQARRQSWFFEKINKIDRPQ
		1				ARLIKKKREKNQIDTIKNDKGD
		ļ				ITTDPTEIQITIREYYKHLYANK
						LENLEEMDKFLDTYTLPRLNQE
						EVESVNRPITGSEIEAITNSLPTK
						KSPGPDGFTAEFYQRYKEELVP
				ŀ		FLLKLFQPIEKEGILPNSFYEASII
						LIPKPGRDTTKKGNFRPISLMNI
		ŀ				DAKIL
19866	50234	Α	19982	1	1698	MELKAKAQELREECRSLRSRC
						NQLEESVSVMEDEMNEMKREG
						KFREKRIKRNEQSLQEIWDYVK
						RPNLRLTGVHESDGENGTKLEN
						TLQDIIQENFPNLARQANIQIQEI
						QRTPQRYSSRRATPRHIIVRFTK
						VEMKEKMLRAAREKGWVTHK
						GKPIRLTVDLSAETLQARREWG
						PIFNILKEKNFQPRISYPAKLSFI
						SEGEIKSFTDKQMLTDFVTTRP
						ALQELLKEALNMERKNRDTHR
						LKIKGWRKIYQANGKQKKAGV
					1	AILVSDKTDFKPTKILKDKEGH
						YIIVKGLIQQEELTILNTYAPSTG
:						APRFIKQVRSDLQRDLDSHTLII
						GDFNTPLSTLDRSTRQKVNKDT
						QELNSALHQADLIDIYRTLHPK
						STEYTFFSAPHHTYSKIDHIVGS
						KALLSKCKRTEIITNYLSDHSAI
						KLELRIKKLTQNHSTTWKLNNL
						LLNDYWVHNEMKAEIKMFFET
						NENKDTTYQNLWDTFKVVCTG
					}	KFIALNAHKRKQERSKIDTLTS
						QLKELEKQEQTHSKASRRQEIT
						KIRAELKEIETQKTL/QKINESRS
		1_				WFFERINKIDRPL
19867	50235	A	19983	1	3450	
19868	50236	В	19984	1	2265	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	l .	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19869	50237	Α	19985	<u>                                     </u>	2526	MFFETNENEDTSYQNLRDTFKV
.,				1	}	VFRWKIIALNAHKRKQERSKID
<u>.</u>	:					TLTSKLKELEKQQQTNSKASRR
						HEITNIRAELKEIETQKTLQKIN
						ESVNWFFEKIIKIDRLLARLIKK
				]		KREKNQIDSIKNDIEDITTDPVE
						TQTTVREYFKHLYANKLENLEE
						MDKFLDTYILPSLSQEEVEPLN
						RPITGSEIEAIINSLPTKKSPGPD
						RFTAKFYQRITNTEKCLKELME/
						PESQSSRTT*RMQKPQESMRST
						GRKGFSDGR*NE*NE*RGEWN
						QVGKHSAGYYPGELPQTSKAG
						QHSDSGNTENATKILLEKSNSK
						TYNCQIHQS*NEGKNVKGSQRE
						RNTNYHQRILQTPLRK*TRKS\K
						EMDKFLDTYTLPRLNEEEVESL
ŀ			i			NTPITGSEIVAIINSLPTKKSPGP
·						DGFTAELYQRQTESQIMSELPF
						TIASKRIKYVGIRLIRDVKDLFK
						ENYKPLLSEIKEDTNKWKNIPC
		1				SWVGRINIVKMAILPKVIYRFN
			· :			AIPIKLPMPFFTELEKTTLNFIW
						NQKRAHIANSILSQKNKAGGIT
						LPDFKLYYKATVTKTAWYWY
						QNREIDEWNTTEPSEIMPHIYN
						YLIFDKPEKNKQWEKDSLFNK
						WCWENWLAICRKLKLDPFLTP
	ŧ					YTKINSRWIKDLNVRPKTIKTLE
						ENLGNTIQDIGIGDDFMSKTPK
	ł					AMATKAKIDRWDLIKLKSFCT
						AKETTIRVNRQPTKWEKIFATY
19870	50238	Α	19986	1	3723	
19871	50239	Α	19987	1	1023	
19872	50240	В	19988	91	1596	
19873	50241	В	19989	1	3171	
19874	50242	Α	19990	1	2952	MLLNQGRKLPRVFAEETLKFK
						GTSNKPQTLEQISTSIIAQKEAT
						VMVPGSNQEIPSGAYAIRALGF
			1			KHKTGRLFEQTLNYLQEFLFTP
		ĺ				QWHLECQQERTVHSPGKAAEA
						REPSVIDRHLIQESSNWHLVGA
}						ALGQSFQRKEQAAIFAVLQPLL
						VIPRQTGSGVDLQKTPTDLQQR
						GLIVRRKTNKQKGIVHNSTTRE
						QNWTENEFDKLTEGGFKRWVI
						TNSSELKEHVLTQCKEDKNLEK
	700:5	ļ			1505	SAIKLELRIKNLIQN
19875	50243	Α	19991	1	1797	

SEQ ID NO:	SEQ ID NO: of peptide sequence	i .	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19876	50244	A	19992		2367	MVKGSIQQEELTNLNMYAPNT GAPRFIKQVLSDLQRDLDSHTLI MGDFNTQLSTLDRSMRQKVNK DTQELNSALHQADLIDIYRTLH PKSTEYTFFSAPHHTYSKIDHIV GSKALLSKCKRTEIITNYISDHS AIKLELRIKNLTQNCSTTWKLN NLLNDYWVHNEMKAEIKMFF ETNENKDTTYQNLWDTFKVVC RGKFIALNAHKRKQERSKTDTL TSQLKELEKQEQTHSKASRRQE ITKIRAELKEIETQKTLQKINESR SWFFERINKIDRPLARLIKKKRE KNQIDAIKNDKGDITTDPTEIQS
						TIREYYKHLYTNKLENLEEMD KFLDTYTLPRLNQEEVESLNRPI TGAEIVAII/NNLPTKKSPGPDGF TAEFYQRK\EGILPNSFYEASIILI PKPDRDTTKKENFRPISLMNID AKILNKILANRIQQHITKLIHHD QVGFIPGMQGWFNIRKSIRVIQ HINRTKDKNYMIISIDAEKAFD KIKQCFMLKTLNKLGIDGTYFK IIRAIYDKPTANIILNGQKLEAFP LKTGTREGCPLSPLLFNIVLEVL ARAIRQEKEIKGIQLGKEEVKLS LFADDMIVYLENPIVSAQNLLK LISNFSKVSGYKINVQKSQAFL YTNNRQTESQIMSELPFTIASKR IKYLGIQLTRDRKDLFKENYKP
19877	50245	A	19993	2	2910	LLKEIKEDTNKWKNIPCSWVEK INIVKMAILPKVIYRFNAIPIKLP WRKIYQANGK/QKKAGVAILVS DKTDFKPTKIKRDKEGHYIMVK
						GSIHQEELTILNIYAPNTGAPRFI KQVLSDLQRDLDSHTIIMGDFN TPLSTLDRSMRQKVNKDTQEF NSALHQADLIDIYRTLHPKSTE YTFFSAPYHTYSKIDHIVQSKAL LSKCKRSEIITNCLSDHSAIKLD LRIKKLTQNHSTTWKLNNLLLN NYWVHNEMKAEIKMFFETNEN KDTTYQNLWDTFKAVCRGIFIA LNAHKRKQETS
19878	50246	Α	19994	1	4729	
19879	50247	Α	19995	1	1566	,
19880	50248	В	19996	1	2796	
19881	50249	В	19997	127	2479	
19882	50250	В	19998	1	2907	

SEQ ID	SEQ ID NO:	1	SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,217	codon for peptide	of peptide sequence	detection, (-possible nucleotide insertion)
			l			
19883	50251	Α	19999	1	3210	MVKGSIQQEELTILNIYAPNTG
			,			ALRFIKQVLRDLQRDLDSHTIIM
						GDFHTPLSTLDRSTRQKVNKDI
						QELNSALHQEDLIDIYRTLHPKS
						TEYTFFSAPHHTYSKIDHIVGSK
		:		1		ALLSKCKRTEIITNCLSDHSAIK
						LELRIKNLTQNRSTTWKLNNLL
						LNDYWVHNEMKAEIKMFFETN
				]		ENKDTTYQNLWDTFKAVCRGK
		]		-		FIALNAHKRKQERSKIDTLTSQL
						KELEKQEQTHSKASRRQEITKIR
		ļ				AELKEIETQ
19884	50252	Α	20000	1	2052	MGDFNTPLSTLDRSMRQKVNK
						DTQELNSALHQADLIDIYRTLH
						PKSTEYTFFSAPHHTYSKIDHIV
						GSKALLSKCKRTEIITNYLSDHS
						AIKLELRIKNLTQNRSTTWKLN
						NLLLNDYWVHNEMKAEIKMFF
						ETNENKDKTYQNLWDAFNAV
						CRGKFVALNAHKRKQEGSKID
						TLTSQLKELEKQEQTHSKASRR
						QEITKVRAELKEIETQKTLQKIN
						ESRS\ARLIKKKREKNQI/DAIKN
						DKRDITIDPTEIQTTIREYYKHL YANKLENLEEMDKFLDTYTLP
				ļ.		RLNQEEVESLNRPITGSEIVAIIN
						SLPTKKSPGPEGFTAEFYQRYK
						EELVPFLLKLFQSTEKEGILPNS
						FYEASIILIPKPGRDTTKKENFRP
						ISLMNIDAKILNKILAKRIQQHI
					·	QKLIHHDQVGFIPGMQGWFNIC
						KSINVIQHINRTKDKNHMIISID
						AEKVFDKIQQRFMLKTLNKLV
İ						LEVLARAVRQEKKIKGIQLGKE
		İ				ELKLSLFADDMIVYLENPIVSA
						QNLLKLISNFSKVSEYKINVQKS
						QAFLYTNNRQTESQIMSELPFTI
				ļ		ASKRIKYLGIQLTRDVKDLFKE
						NYKPLLKEIKEDVNKWKNIPCS
						WVGRINIVKMAILPKVIYRFNAI
						PIKLPMTSFTELEKTTFKFIWNH
						KGARIAKSILSQKNKAGGITLPD
						FKLYYKATVTKTAWY
			<u> </u>	L	<u> </u>	TAREL LINES A LINES AND I

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
ŀ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19885	50253	Α	20001	1	2314	MDKFLDTYTLPRLNQEEVESLN
						RSITGSEIVAILNSLPTKKSPGPD
						GFTAEFYQRYKEELVPFLLKLF
			1			QSIEKEGILPNSFYEASIILIPKPG
						RDTSKKENFRPIPLMNIDAKILN
						KILAKRIQQHIKKLIHHDQVGFI
						PGMQGWFNIRKSINVIQHTNRA
						KDKNHMIISIDAEKAFDKIQQLF
						MLKTLNKLGIDGMYLKIRRAIY
1						DKPTANIILNGQKLEAFPLKTGT
						RQGCPLSPLLFNIVLEVLARAIR
						QEKEIKGIQLGKEEVKLSLFAD
						DMIVYLENPIVSAQNLLKLISNF
						SKVSGYKINVQKSQAFLYTNNR
1						QTESQIMSELPFTLASKRIKYLG
1						IQLTRDVKDLFKENYKPLLNEI
						KEDTNKWKNIPCSWVGRINIVK
		1				MAILPKVIYRLNAIPIKLPMPFF
						TELEKTTLKFIWNQKRARVAKS
ļ						ILSQKNKAGGITLPDFKLYYKAI
ļ						VTKTAWYWYQNRDIDQWNRT
						EPSEITPHINNYLIFDKPDKNKQ
						WGKDSLFNKWCWENGLAICRK
						LKLDPFLTPYTKINSRWIKDLN
						VRPKTRKTLEENLGITIQDIGMG
		l				KDFMSKTPKAMATKDKIDKW
						DLIKLKSFCTAKETTIRVNRQPT
						KWEKIFATYSSDKGLISRIYNEL
						KQIYKKKTNNPIKKWAKDMNR
						HFSKEDIYAAKKHMKKCSSSLA
						IREMQIKTTMRYHLTPVRMAII
	}					KKSGNN\RLFTVAKTWIQPKCA

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19886	50254	Α	20002	1	2685	MGDFNTPLSTLDRSTRQKVNK
						DTQELNSAPHQADLIDIYRTLH
						PKSTEYTFFSAPHHTYSKTDHIL
						GSKALLSECKRTEIITNYLSDDS
		1				AIKLELRIKNLTQNRSTTWKLN
! :						NLLLDDYWVHNEMKAEIKMFF
ŀ						ETNENKDTTYQNLWDAFKAVC
						RGKFVALNAHKRKQGRSKIDT
						LTSQLKELEKQEQTHSKASRRQ
						EITKIRAELKEIETQKTVQKINES
ŀ			!			RSWFFERINKIDRQLARLIKKKR
	:					EKNLIDAIKNDKGDITTDPTEIQ
						TTIREYYKHLYANKLENLEEM
						DKFLDTYTLPRLNQEEVESLNR
						PITGSEIVAIINSLTTKKSPGPDG
1						FTAEFYQRAIRQEKEIKGIQLGK
						EEVKLSLFADDMIVYLENPIVS
						AQKLISNFSKVSGYKINVQKSQ
	,					AFLYTNNRQTESQIMSELPFTIA
		l				SKRIKYLGIQLTRDVKDLFKEN
		l				YKPLLKEIKEDTNKWKNIPCSW
		1				VGRINIVKMAILPKVIYRFNAIPI
		1		:		KLPMTFFTELKKTTLNFIWNQK
		l		į		\RAHIAKS/VLSQKNKAGGITLP
		1				DFKLYYKATVTKTAWYWYQN
	1	1				RDTDQWNRTEPSEIMPRIYNYL
				1		IFDKPEKNKQWGKDSLFNKWC
						WKNWLAICRKLKLDPFLTPYT
						KINSRWIKDLNIRPKTIKTLEEN
						LGITIQDIGMGKDFMSKTPKAM
						ATKAKIDKWDLIKLKSFCTAKE
						TTNRVNRQPTKWEKIFATYSSD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	I .	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		:
19887	50255	Α	20003	1	2502	MTELTGIQQPQIVLFEHKGHKL
						VQGSSSDAGKVNRIYQHYEAS
				l		DKFNYTTGLAWKTAPEQTGKT
			1			VRKQQIKLNVKKMESRSKMQE
						HSSSPPMEQSWRENDFDELREE
						AFRRSNYSELQEEIQTKGQEVK
						NFEKTLDEYITRITNTEKCLKEL
						MELKAKARELREECRSLRSRCD
ļ						QLEERVSVMEDEMNEMKREG
						KFREKRIKRNEQSLQEKWDYV
						KTPNLRLIGVPESDGENGTKLE
						NTLQDIIQENLPNLVRQANIQIQ
						EIQRTPQRYSSRRATPRHIIVRFT
				•		KVEMKEKMLRAAREKEIQTTIR
						EYYKHLYANKLENLEEMDKFL
						DTYTLPRLNQEEVESLNRPITGS
						EIVAIINSLPTKKSP/GPVGFTAE
						FCQRK\EGILSISFCEASIILIPKL
						GRDTTKKENFRPISLMTIDTKIF
						NKILANQIQQHIKKLIHHDQVG
						FIPGMQGWFNICKSINVIQHINR
				•		TKDKNHMIISIDAEKAFDKIQQL
						FMLKTLNKLGIDGTYFKIIRAIY
						DKPTANIILNGQKLEAFPLKTGT
						RQGCPLSPLLFNIVLEVLAGAIR
						QEKEIKGVQLGKEEVKLSLFAD
						DMIVYLENHIVSAQNLLKLISNF
						SKVSGYKINVQKSLAFLYTNNR
						QTESQIMSELPFTIASKRIKYLGI
						QLTRDVKDLFKENYKPLLNEIK
						EDTNKWKNIPCSWVGRINIVK
						MAILSKVIYRFNAIPINLPITVFT
19888	50256	В	20004	1	2115	
19889	50257	Α	20005	1	2382	
19890	50258	Α	20006	1	2742	
19891	50259	A	20007	1	2250	
19892	50260	Α	20008	1	2982	
19893	50261	Α	20009	1	4791	
19894	50262	Α	20010	1	2499	
19895	50263	Α	20011	1	3259	
19896	50264	Α	20012	1	3654	
19897	50265	Α	20013	1	2391	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
]				sequence		
19898	50266	Α	20014	326	1158	LWICEVMVKQRSRILKICLQKQ
						WQKTLAMWLKPCMGTFLLQL
					Ì	C*LDIAWV/RAIAVHTASSNLVP
						SLLGLCMIDVVEGTAMDALNS
						MQNFLRGRPKTFKSLENAIEWS
		ŀ				VKSGQIRNLESARVSMVGQVK
						QCEGITSPEGSKSIVEGIIEEEEE
		İ				DEEGSESISKRKKEDDMETKKD
	•					HPYTWRIELAKTEKYWDGWFR
						GLSNLFLSCPIPKLLLLAGVDRL
					[	DKDLTIGQMQGKFQMQVLPQC
						GHAVHEDAPDKVAEAVATFLI
						RHRFAEPIGGFQCVFPGC
19899	50267	A	20015	1	2860	
19900	50268	Α	20016	1	2956	
19901	50269	Α	20017	1	3359	MNNSIKEKVKLSLTKWEMQTK
						PNNNTVDLKDAQEVNLTSTDY
						RKQHVMLPTEINSDDSLLKIGVI
						RLANWIKSQDPSVCCNQETHLT
						CRDTHRLKIKGWRKIYQANGK
						QKKAGVAILVSDKTDFKPTKIK
			1	1		RDKEGHYIMVKGSIQQEELTIL
1						NTYAPNTGAPRFIKQVLSDLQR
						DLDSYTLIMGDFNTPLSTLDRS
ł						TRQKVNKDTQELNSALHQVDL
						IDIYRTLHPKSTEYTFFSAPHHT
		<u> </u>				YSKIDHILGSKALLS
19902	50270	A	20018	3	3130	
19903	50271	В	20019	8	2398	SNSHITILTLNVNGLNAPIKRHR
19904	50272	Α	20020	3	3398	1
						LANWIK\SQDP\SVSCIQETHLTC
						RDTHRLKIKGWRKIYQANGKQ
						KKST\VAILVSDKTDFKPTKIKR
						DKEGHYIMVKG*IQ*EELTILNI
						YAPNTGAPRFIKQVLRDLQRDL
						DSHPIIMGDFNTPLSTLDRSMR
						QKVNKDIQDLNSALHQADLTDI YRTLHPKSTEYTFFSPPHHTYSK
	-					TDHIVGSKALVSKCKRTEIITNC
						LSDHSAIKLELRIKKLTQNRSTT
10005	50272	D.	20021	102	2210	WKLNNL
19905	50273	В	20021	483	12210	<u> </u>

SEQ ID NO:	SEQ ID NO: of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19906	50274	A	20022		5316	MVKGSIHQEELTILNIYAPNTG APRFIKQVLSDLQRDLDFHTLI MGDFNTPLSTMDRSTRQKVNK DIQELNSALHQADLIDIYRTLHP KSTEYTFYSAPHHTYSKIDHILG SKALPSKCKRTDIITNYLSDHSA IKLELRIKNLTQNHSTTWKLNN LLLNDYWVHNEMKAEIKMFFE TNENKDTTYENLWDAFKAVCR GKFIALNAHKRKQERSKIDTLT SQLKELEKQEQTHSKASRRQEI TKIRAEQINKIDR
19907	50275	A	20023		2712	WRKIYQANGKQKKAGVAILVS DKTDFKPTKIKRDKEGHYIMVK GSIQQEELTILNIYAPNTGAPTFI QQVLSDLQRDLDSHTLIIGDFN TPLSTSDRSTRQKVNKDTQELN SALHQADLIDIYRTLHPKSTEYT FFSAPHHTYSKIDHILGSKALLS KCKRTEIITNYLSDHSAIKLELM IKNLTQNHSTIWKLNNLLLNDY WVHKEMKAEIKMFFETNENKD TTYQNLWDTFKAVCRGKFIAL NAHKRKQERSKIDTLTSQLKEL EKQEQTLSKASRRQEITKIRAEL KEIETQKTLQKINESRSRFFERIN KIDR*\TLARVIKKKGEKNQIDTI KNDKGDITTDPTEIQTTIRESYK HLYANKLENLEEMDKFLDTYT LPRLNQEEVESLNRPITGSEIVAI VNSLPTKKSPGPDGFTAEFYQR YKEELVPFLLKLLQSIEKEGILP HSFYEASIILIPKPGRDTTKK\EN FRPISLMNIDAKILNKILANQIQ QHIKKLIHHDQVGFIPGMQGWF NTCKSINVIQHINRTKDKNHMII SIDAEKAFDKIQQFFMLKTLNK LVLEVLARAIRQEKEIKGIQLGK EEVKLSLFADDMIVDLENPIVS AQNLLKLISNFSKVSGYKINVQ KSQAFLYNNNRQTESQIMSELA FTIASKRIKYLGIQLTRDVKDLF NENYKPLLNEIQEDTNKWKNIP CSWVGRINIVKMAILPKVIYRF

SEQ ID		ŧ .	SEQ ID NO:	Į.	l .	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
ţ	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19908	50276	Α	20024	1	4461	MVASRSSRVGVFYGGQNSVDG
Į.	ļ		ļ	į		VSRRSDGSLNELCLYRPPTGAY
						LAYDTLDVLPSGQFTVNTPPID
	1					VNGKSLLALLWEHTSLLTSMG
						GGRWAMQPLGQLGQLEHLGD
}						RRDQVLDRSGRSRRLDAAQHL
						RHRPDRRPQGMVGVLTNQKEP
						RTRWIHSRILPEVQGGTGTVPSE
į.						TIPIDRKEGILPNLFDEASIILIPK
						RGRDTTKKENFRPISLMNIDAKI
						LNKILANRNQQHIKKLIHHDQV
						GFIPGMQGWFNICK
19909	50277	В	20025	1183	4247	
19910	50278	A	20026	1	2671	MVKGSIQQEELTILNIYAPNTG
				_		APRFIKQVLSDLQRDLDSHTLI
		ļ	ļ			MGDFNTPLSTLDRSTROKVNK
						DTQELNSALHQADLIDINRTLH
						PKSTEYTFFSAPHHTYSKIDHIV
				÷		GSKALLSKCKRTEIITNYLSDHS
						AIKLELRIKNLTQSRSTPWKLN
						NLLLNDYWVHNEMKAEIKMFF
						ETNKNKDTTYQNLWDAFKAV
						CRGKFIALNAYKRKQERSKIDT
						LTSQLKELEKQEQTHSKASRRQ
		ĺ '				EITKIRAELKEIETQKTLQKINES
			l i			RSWFFERINKIDRPLARLIKKKR
						EKNQIDTIKNDKGDITTDPTEIQ
						TTIREYYKHLYANKLENLEEM
						DTFLDTYTLPRLNQEEVESLNR
		1				PITGSEIVAIINSLPTKKSPGPDG
						FTAEFY/PESYL*QTHSQYHTEW
						AKTGSIPFENWHKTGMPSLTAP
				!		IQHSVGSSGQGNQAGEGNKGY
						SIRKRGSQIVPVCR*HDCLSRKP
					(	HRLSPKSP*ADKQLQQSLRIQN
				i		QCTKITSILIHQKQTNREPTHE*
						TPIHNCFKENKIPRNPTYKGCEG
						PLQGELQTTAQRNKRGHKQME
		(				EHSMLMGRKNQYRENGHTAQ
						GNLQIQCHPLQATNDFLHRIGK
				,		NYFKVHMEPKKSPHRQVNPKP
				,		KEQSWRHHTT*LQTILQGYSNQ
						NS\MVLVPKQRYRSMEQNRAL
						RNNAAYLHYSD\L\*QT*EKQA
						WGKGIP\IYKM\AL\GKLAS\PM*

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19911	50279	Α	20027	1	3159	MGDFNTPLSTLDRSSRQKVNK
						DTQELNSTLHHADLIDIYRTLHP
						KSTEYTFFSAPHHTYSKIDHVV
						GSKALLSKCKRTEIITNCLSDHS
	•					AIKPELRIKKLTQNRSTTWKLN
						NLLLNDYWVHNKMKAEIKMFF
						ETNENKDTTYQNLWDTFKAVS
						RGKFIALNAHKRKQKRCKIDTL
						ASQLKEVEKQEQTHSKASRRQ
		İ		ļ		EITKIRAELKEIETQKTLQ\KISES
						WSWFFEKINKIDRPQARLIKKK
						REKNQIDTIKND
19912	50280	Α	20028	2	3767	HRLANWIKSQDPSVCCIQEAYL
						TCRDTHRLKIKGWRKIYQANG
						KQKKAGVAILVSDKTDFKPTKI
						KRDKEGLYIMVKGSIPREELTIL
						NIYAPNPEAPRFTKQVLSDLQR
j						DVDSHTIILGDFNTPLSALDRSM
						RQKVNKDTQELNSALHQADLI
	:					DIYRTLHPKSTEYTFFSAPHHTY
				ļ		SKIDHRVGSKALLSKCKRSEIIT
						NCLSDHSAIKLELRIKKLTQNCS
						TTWKLNNLLLNDYWVHNEMK
j						AEIKMFFETNEN
19913	50281	Α	20029	3311	9648	KLYVCITRITNAEKYLKELMEL
						KAKAQELCEECRSLRSRHDQLE
						ERVSVMEDQMNEMKQEEKFRE
						KRRKRNEQSLQEIWDYVKRPN
		1				LRLIGVPESDGENGTKLENTLQ
						DIIQKNFPNLARQANIYIQEIQEI
			:			QRTPQRYSPRRATPRHIIVRFTK
			]			VDMKEKMLRAVREKGFKPTKI
						RRDKEGHYIMVKGSIQQEELTI
						LNIYAPNTGGPRFIKQVLSDLQ
						RDLDSHTIIMGDFNTPLSTLDRS
			l			MRQKINKDIQEL
19914	50282	Α	20030	2031	6332	QDSNSHITILTLNI\NGLNAPI*R
						HD\RNCNWIKSQDPSVCCIQET
[						HLTCRDTHRLKIKGWRKIYQA
						NGKQKKKAGVAI\LVSDKTDFN
						PTKIKRDKEGHYIMVKGSIQQE
			İ	1		ELTILNIYAP\NTGAPRFIKQVLS
			1	l .		DLQRDLDSHTLI\MGD\FNTPLS
				1		TLDRSRRQKVNKDTQELNSAL
						HQADL\IDIYRTLHP\KSTEYTFF
						SAPHHTYSKIDHIVGSKALLSK
				}		CKRTGIITNYLSDHSAIKLELRI
						KNLTQSR
19915	50283	Α	20031	2	178	
19916	50284	С	20032	88	228	

SEQ ID	SEQ ID NO:				1	Amino acid sequence ( X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		07/340,217	sequence	or peptide sequence	detection, (~possible nucleotide insertion)
				•		
19917	50285	Α	20033	3	382	VDRTIMSDSLVVCEVDPELTEK
		l				LRKFRFRKETDNAAIIMKVDKD
	İ					RQMVVLEEEFQ\FVVYSYKYV
						HDDGRVSYPLCFIFSSPVGCKPE
						QQMMYAGSKTRLVQTAELTK
			_			VQTGMWLQSVRERWCGSWV
19918	50286	Α	20034	89	633	KTAELRKEEACGQNNHALPGS
						LRQSDSLVVCEVDPELTEKLRK
						FRFRKETDNAAIIMKVDKD/RG
						QMVVLEEEF\QNISPQREL\KMG
						FPRRDKPR\FVVYS\YKYVHD\D
						G\RVSYP\LCFIFPPRPVGCKPE\Q
			ŀ			QMMY\AGS\KNRLV\QTAELTK
						GVPKSRTT*LTSLRPWL\QEKLS
						FLSLISGAGD
19919	50287	Α	20035	78	638	WLDLTWKESSADSQLFLEKMD
						HSHHMGMSYMDSNSTMQPSH
						HHPTTSASHSHGGGDSSMMM
						MPMTFYFGFKNVELLFSGLVIN
				1		TAGEMAGAFVAVFLL\AMFYE
						GLKIARESLLRKSQVSIRYNSMP
						VP\GQNG\TILYGRTHKTVG\QQ\
						MLSFPHLLQTVLHIIQV\VISYFL
	<u> </u>					MLIFMTYNGYLCIAVA
19920	50288	A	20036	1	849	MSQKMENLIGNAHISAYEIIKEF
		1		}		QKEQHQAGLPGTTVPCSCQVY
		1				KCSKRPERVAREGPSEDLEPLP
		1				QTVRKCGQGCTLHGAGGSQGQ
		1				VGARPFGIGGVEALWFLASPSY
			1			WAPLIPQCQLGKLLVVRLVQL
		l	-			QARRELAPMLAPGAACPMVAT
						ACLTVQWPDPMLTHTPLTIPHL
		l				TCSLLGGNVELLFSSLVINTAGE
		l				MAGAFVAVIILAVFYEGL\IAQE SLL*PNGTILM*THKTVGQQML
		1				SFPHLLQTVLHIIQVVISYLLMLI
						FMTYNGSSELLFRGERPR
19921	50289	В	20037	308	568	FMI INGSELLFRGERFR
19922	50290	A	20038	394	528	
19923	50291		20039	186	856	TWGKGDILRRPRRKMSSY/SHF
						FVQTCRGG\HKKKHPRLLQVN\
						FSEF\SKEVLQRGWRPMSCLKR
						KGKILKIWAKAGQGPFLERELK
						TLYPLPKGEDQKKKFKDPNCTP
						RGPPS\AFFPLLALEYPPQKSKG
			:			EHPWPVPLVDVAEENWETMW
						ELTLLQMDKQP\YEKKA\AKLK
						EKYEKDIACISSLKGKPDAAKK
						RELSRPEKSKKKKEEEEDEEDE
						EDEEEEEDE*DEEDEEEEDE
L	i	Ц	<u> </u>		l	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19924	50292	Α	20040	1	413	GAMRGDRGRGRGGRFGSRGGP
		ŀ				GGGFRPFVPHIPFDFYLCEMAFP
				ĺ		RVKPAPDETSFSEALLKRNQDL
		ľ				APNSAEQASILSLVTKINNGNE
						KL\VGSPGTLKAHIEKFTRGIPY
						KRGTVSRGHNVADLGGKLKIL\
						PPLEAV
19925	50293	A	20041	70	1451	GAMRGDRGRGRGGRFGSRGGP
						GGGFKPFVPHIPSDFYLCEMAF\
	l					PRVKPAPDETSFS*GLA*RRNQ
		İ				D\LAPNSAEQ\ASILSL\VTKI\NN
						V\IDNLTCGSQGHFEVAKLEEV\
						RQVGDPIKRGTMTTGHNVA*/L
						WVVIPQEFWQRLEA\VACPGGN
						KVVESLR/SHQDPSESFKPCLTN
						E\TGFEISSSDATVKILITT\VPP\N
	ŀ	l				LRKLDPELHLDIKVLQSALAAI
						RHARWFEENASQSTVKVLIRLL
	ŀ	l				KDLRIRFPG\FEPLTPW\ILDLLG\
		İ				HY\AVMNNPTRQPLGLKRWHN
						RALAWQISGLAGTVSCQG\SVG
	}					YSIDPCESWQL*ELHTVMTLEQ
		l				Q\DMV\CYTAQ\TFVRILP\HGGF
						RKILG\QEVDASYLASEIST\WD
		l				GVIVTPFRKRLYEKP\PEK\KEG
		1				EEEEENTEEPTSRRGRRK\HGN\
		1				SGVTIPLPLLFPNPRGKTGALSC
						PAYWALHGDRHFPWDREDSRK
19926	50294	Α	20042	253	379	
19927	50295	Α	20043	284	1062	APRNYLPQRATASVRRQQISFS
						NEHTPHQSLHEIPNVRQRHSGL
						MKLNTMGAMTTSTTSAVANH*
						PSNKRTPLVSV/MQSASPEQRQ
						AVRERMQTNPKIQQRREAARE
						RIQSASPEQRQAVREKMQTNPQ
						NQQRRDAARERIQSASPEQRQV
						FKEKVQQRPLNQQQRDNARQR
						VQSASPEQRQVFREKVQESRPQ
						RLNDSNHTVRLNNEQRSAVCE
						RLSERGARRLESLYGRHMQAN
						PEPPKKNNDKSKKISRKPLAAK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				Sequence		
19928	50296	Α	20044	1	2269	MSPDIPQVSPEVGLYPVANHSSI
						VVPTCHSLSEEDTEAQRNHELD
		ł				TVKALREDAQVLKIPNQSPKDT
						LVTALQLAQKGTKAAAIRLGA
	-					KVGQKQSRQAKSSPALVVPGL
					:	LSFPRARFFEWQSPGLPNTAAM
						ENGTGPYGEERPREVQETTVTE
		İ				GAAKIAFPSANEVFYNPVQEFN
						RDLTCAVITEFARIQLGAKGIQI
						KVPGEKDTQKVVVDLSEQEEE
						KVELKESENLASGDQPRTAAV
		İ				GEICEEGLHVLEGLAASGLRSIR
						FALEVPGLRSVVANDASTRAV
						DLIRRNVQLNDVAHLVQPSQA
		1				DARMLMYQHQRVSERFDVIDL
						DPYGSPATFLDAAVQAVSEGG
			:			LLCVTCTDMAVLAGNSGETCY
				1		SKYGAMALKSRACHEMALRIV
						LHSLDLRANCYQRFVVPLLSIS
1		İ				ADFYVRVFVRVFTGQAKVKAS
						ASKQALVFQCVGCGAFHLQRL
		-				GKASGVPSGRAKFSAACGPPVT
						PECEHCGQRHQLGGPMWAEPI
						HDLDFVGRVLEAVSANPGRFH
		1				TSERIRGVLSVITEELPDVPLYY
		ł				TLDQLSSTIHCNTPSLLQLRSAL
						LHADFRVSLSHACKNAVKTDA
İ			1			PASALWDIMRCWEKECPVKRE
						RLSETSPAFRI\LSVEPRLQANFT
						\IRED\ANPSSRQRGLKRFQA\NP
						EA\NWGPRPRSRP\GGKAADEA
		ļ			0.000	MEER\RRLLQNKRKEPPEDV\A
19929	50297	Α	20045	1578	2356	PLRRWACNHKLRQNAHQLSSTI
						HCNTPSLLQLR\SALLHADFR\V
						SLSHACK\NAVKTDAPASALW\
						DIMRCWEKECPVKRERLSETSP
						AFRILSVEPRLQANFT\IREDANP
1						SSRQRGLKRFQANPEANWGPR
1						PRARP/GVSGGKVGVGLSQGKK
1			1			G*RADGVPPTPSPSHRGKAADE
1			1			AMEERRILLQNKRKEPP\EDVA
						QRAARLKTFPCKRFKEGTCQRG
1						DQCCYSHSPPTPRVSADAAPDC
L	L	<u> </u>	L	l	<u> </u>	PETSNQTPPGPGAAAGPGID

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19930	50298	A	20046		791	MGPSSIHHSSQHATAPQILPKVR AGPRSSGCRISECRREEKPPKGR RGLKDATRPGVNWGPDGGVQS GAGAQPWSLTPLPPGSVSPAGR AQRRSMRAAAAQERSPGRRRL GKEPLIRESWTRSAPWRCLSQD AGRAGSQVCGSSPQTRGAAAE KVAVTGAKSCSGRGKKAAAAR AKSRGGGGRKTQKAAAAKSRG C/AGQRSPKSRGSRG*IPRGRGQ KAAVAGAESNGGGGKKSRRRR HKAAKTRSSGGKNPWRQKAAS AGAK
19931	50299	С	20047	76	327	
19932	50300	A	20048	336	814	ANFSLEDPSCSPPPRLLAGSFVD SPQSLSCMRSKNPLLGSGSGPLS PTYIL*ATTPHFLLPQPLITTSPSS PSPPSPPLPPPPPSFSSSYSSSSFF E/NRVPLCCPCFGSAIRKPSPSVS TKARPQGETGRSLALLFLPQRP SSSLHRSDTLYRDCVHRI
19933	50301	Α	20049	1	310	
19934	50302	Α	20050	1	1482	
19935	50303	Α	20051	3	2455	
19936	50304	A	20052	3	1441	

2534

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19937	50305	A	20053		2322	MAEGKGEAGTFFTGQQDRLLF VFFIITIVVGLFRCSVFSSSSSSS LSSSSSSSSSSSSSSSLHLFFIFIF VFFVFIFFVFIFVFVFAFFFIFFVF VVFFVFFFFVLLVEILMRPTISIR GQKLKISDEMSKDCLSILYNTC VCTEGVTKRLAEKNDFVIFLFT LMTSKKTFLQTATLIEDILGVK KEMIRLDEVPNLSSLVSNFDQQ QLANFCRILAVTISEMDTGNDD KHTLLAKNAQQKKSLSLGPSA AEINQAALLSIPGFVERLCKLAT RKVSESTGTASFLQELEEWYT WLDNALVLDALMRVANEESEH NQASIVFPPPGASEENG\LAHTS
						ART\QLPQSMKIMHEIMYKLEV LYVLCVLLMGRQRNQVHRMIA EFKLIP\G\LNNLFDKLIWRKHS ASALVLHGHNHY\CDCSPDITL\ KIQFLRLLQSFSDHHENKYLLL NNQELNELSAISLKANIPEVEAV LNTDRSLVCDGKRGLLTRLLQ VMKKEPAESSFRFWQARAVEE FPSRGTTSYADQMFLLKR\GLL
						EHILYCIVDS\ECKSRDVLQSYF DL\LGEL\MKFNVDAFKRFNKYI NTDAKFQVFLK\QI\NSSLVDSN MLVRCVTLSLDRFENQVDMKV AEVLSECRLLAYISQ\VPTQMSF LFRLINIIHVQTLTQENVSCLNT\ SLVILML\ARRKERLPL\YLRLL\ QR\MEHSKKYPGLL\LNNFHNL

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19938	50306	Α	20054	237	2033	TSPTPLPLPATMPKTISVRVTTM
						DAEL\EFAIQPNTTGKQLFDQV
						VKTIWLEGKFGFFGLQNQD1\K
						GFST\WLKLNRKVD*PRNVRKE
-						\SPLLFK\FRAKFYP\EDVSEELIQ
						DITQRLF\FLQVKEGILNDDIYCP
						\PETAVLLASYAVQSKYGDFNK
						EVHKSGYLAGDKLLPQRVLEQ
į						HKLNRD\QWEERIQVW\HEEHR
						GML/RGEDAVLEISERLLQDLE
						MYGVNYFSI\KNKKGSELWLG
ļ						VDALGLNIYEQNDRLTPKIGFP
}						W\SEIRNISFNDKKF\VIKPIDKK
1						APDFVFYAPRLRINKRILALCM
						GNHELYMRRRKPDTIEVQQMK
						AQAREEKHQKQMERAMLENE
						RKKREMAEKEKEKIEREKEEL
						MERLKQIEEQTKKAQQEL\EEQ
						TPRAPGTLSQERKRAQSEAEKL
ŀ						AKERQEAEEAKEALLQASRDQ
						KKTQEQLALEMAELTARISQLE
						MARQKKESEAVEWQQKAQMV
						QEDLEKTRAELKTAMSTPHVA
						EPAENEQDEQDENGAEAS\ADL
						RA*S\MAKDRSEEERTTE\AEKN
						ERVQKHLKALTSELANARDES
						KKTANDMIHAENMRLGRDKY
						KTLRQIRQGNTKQRIDEFESM

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19939	50307	A	20055	2	2230	PRVRLGAHGCPPGPRRCGCCW ANGLRSSERCVLESLRPCVQSA
					ļ	TRGLRCGCDSAANLDTMAKPP
İ						QGAGDRGSIHDFPGFDPNQDAE
						ALYTAMKGFGSDKEAILDIITSR
ŀ						SNRQRQEGCHSYK\SLYGK\DLI
		1				ADLK\YELTGKF\ERLIVRLMRP
						PAYCDAKEIK\DAISG\IGTDEK\
						CLIEILASRTNEQMHQLVKRITK
ŀ						MPTERDLEADIIGDTSGHFQKM
						LVVLLQGTREEDDVVSEDLVQ
1						QDVQDLYEAGELKWGTDEAQF
ŀ						IYILGNRSKQHLRLVFDEYLKT
						TGKPIEASIRGELSGDFEKLMLA
						VVKCIRSTPEYFAERLFKAMKG
						LG\TRDNTLIRIMVSRSELDMLD
		]				IREIFRTKY*/EKSLYSMD/IKND
						TSG/QSYKKTLLKLSGGDDDAA
1						DQFFPKAAQVAYQMWELSAV
		İ				ARVELKGTV\RPANDF\NP\DAD
						AKALR\KAMKGLGTDEDTIIDII
						THRSNVQRQQIRQTFKSHFGRD
						LMTDLKSEISGDLARLILGLMM
						PPAHYDAKQLKKAMEGAGTDE KALIE\ILATRTNAEIRAINEAYK
						\EDYHKSLEDALSSDTSGHFRRI
	1					LISLATGHREEGGENLDQARED
						AQVAAEILEIADTPSGDKTSLET
						RFMTILCTR\SYPHVRRVIQEFIR
						LTNYDVEHTIKK\EM\SGDIRDA
						F\VAIVQ\SAKTKPLFFPDK\LYK
		ŀ				SMKG\AGTDEKTLTRIM\VSR\G
19940	50308	Α	20056	3	368	
19941	50309	A	20057	1	936	MMKCEEQKEEEEETVRMFSAA
	1					SOPLDPDGTLFRLRFTAMVWW
İ		ľ				AITFPVFGFFFCIIWSLVFHFEYT
						VATDCGVSASSPAPGSGPG\PN
						YLPSVSSAIGGE\VPQRYVWRFC
						IGLHSAPRFLVA\FAYWNHYLS
						CTSPCSCYRPLCRLNFGLNVVE
						NLALLVLTYVSSSEDFTIHENAF
						IVFIASSLGHMLLTCILWRLTKK
						HTVSQEVRSIPSGGSKAAQKKI
						KDICPQDSGNGEDRKSYSWKQ
						RLFIINFISFFSALAVYFRHNMY
						CEAGVYTIFAILEYTVVLTNMA
						FHMTAWWDFGNKELLITSQPE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19942	50310	Α	20058	119	930	ESLFPSLAFLSPSKPLFSHPPSFH
						LSNKQDTSPCSCVLSSPFSLRTE
						ASIGVPGIKMDGGNAQEHGHFL
						RLPQVPNYLPSVSSAIAREVPQR
						YVWRFCIGLHSAPRFLVAFAY
						WNHYLSCTSPCSCYRPLCRLNF
						GLNVVENLALLVLTYVSSSEDF
						/TWVPG*GRSGEVFPEGTGLPLP
						HSDLPTSWCGHSLQCGSQSSFP
						PAIHENAFIVFIASSLGHMLLTCI
						LWRLTKKHTVSQEVRSIPSGGS
						KAAQKKIKDICPQDSGNGEVGT
						EF
19943	50311	Α	20059	3	206	
19944	50312	В	20060	80	304	
19945	50313	Α	20061	1	2111	MAATGTAAAAATGRLLLLLV
	1					GLTAPALALAGYIEALAANAGT
						GFAVAEPQIAMFCGKLNMHVN
						IQTGKWEPDPTGTKSCFETKEE
	•					VLQYCQEMYPELQITNVMEAN
	1					QRVSIDNWCRRDKKQCKSRFV
	1					TPFKCLVGEFVSDVLLVPEKCQ
	-					FFHKERMEVCENHQHWHTVV
						KEACLTQGMTLYSYGMLLPCG
			}			VDQFHGTEYVCCPQTKIIGSVS
ļ						KEEEEEDEEEEEEDEEEDYDV
	1					YKSEFPTEADLEDFTEAAVDED
						DEDEEEGEEVVEDRDYYYDTF
						KGDDYNEENPTEPGSDGTMSD
						KEITHDVKVPPTPLPTNDVDVY
						FETSADDNEHARFQKAKEQLEI
						RHRNRMDRVKKEWEEAELQA
•		i				KNLPKAERQTLIQHFQAMVKA
ļ						LEKEAASEKQQLVETHLARVE
						AMLNDRRRMALENYLAALQS
						DPPRPHRILQALRRYVRAENKD
						RLHTIRHYQHVLAVDPEKAA\Q
						MK\SQVMTHL\HVIEERRNQSLS
						LLYKVPYVA\QE\IQEEIDDVLQ
						EQRADMDQFTASISE\TPVDVR
						V\SSDESEEIPP\FHPFHPFPAPT*
						ERRILGVGRAGWGGLIGAEEK
				1		VINSK\NKVDE\NMVH*TRLLD
	1	1				VKEMIFNAERVGG\LEEERESV
						GPLREDFSL\SSSALIGLLVIAVA
						IAT\VIVISLVMLRKRQYGTI\SH
		<u> </u>		<u></u>		GIVE/VVDPMLTPEERHLNKMQ

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19946	50314	A	20062	61	2634	SQPAEGKWSTGNAATATAGVL LLPAACGA*TAPALAL\AGYIEA LAANAGTGFAVAEPSNSPMFC GKLNMPCETFQTGK\WEP\DPN QAPKSCF*NQKEEVSSSYF\QEM YPRAYRSQNVMEAKPAGLRID N\WC\RRDKKQ\CKSRFCYTFSS CLVG*IL*SDVPAKFPEKVPSFF HK\ERMEVCE\NHQ\HW\HTVSQ KKAMS*PPGK*PLIIANGMPASP CGGRTSFHGTE\YVC*PSAQRLL GSCV/TKEEEEEEE\EKEEEEDE KGD\YAVYKGNFLL*ADLEDST EA\AVDEDDEDEREGEDVVEDR EYYYDTFKGDDYNEENPN\DPG SAATMSD\KEITHD\VKAVCSQE A\MTGPCRAVMPLWYFYLSKG KCVRFIYGGCGGNRNNF*SEDY CMAVCKAMIPPTPLSTNDDDVF F*TSKDDNEHARFHKAKEHVVI RPRNRMDRLK*EW*YAQL\QLK \NLPKAERQ\TLIQHF\QA\MVKA LEKEAA\SEKQQLVET/HTLARV EAMLND\RCLMALENY\LAALQ SDP\PPHR\ILQALRRYVRAEN KDRLHTIRHYQHVLAVDPEKA AQM\KSQVM\THL\HVIEERRNQ SLSLLYKVPYVAQEIQEEIDELL QEQRADMDQ\FTASISE\TPVDV \RVS\SEGRGEIPPFHPFHPFPALP
19947	50315	A	20063	734	1049	ENEDTQPV\LYHPMKKGSGVGE QDGGLIVAEEK\VINSK\NKVDE CCPPGGTGRCLGTCGYHN/CGC SI*SSDLPQR/CCVSGRR*TFWM
10048	50216		20064	6	1174	VWSACVCASGCLPGGNRCCCP VSLSPERIAVLGGLLPRKPPCRV VGERRSVLTLQATSQCPR
19948	50316	A	20064	285	567	SRRRSSSYERSGSYSGRSPSPYG RRRSSSPFLSKRSLSRSPLPSRKS MKSRSRSPAYSRHSSSHSKKKR SSSRSRHSSISPVRLPLNSSLGAE LSRKKKERAAAAAAAKMDGK ESSYERSGSYSGRSPSPYGRRRS SSPFLSKRSLSRSPLPSRKSMKS RSRSPAYSRHSSSHSKKKRSSSR SRHSSISPVRLPLNSSLGAELSR KKKERAAAAAAKMDGKESK GSPVFLPRKENSSVEAKDSGLE SKKLPRSV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19950	50318	A	20066	214	805	LGLQAANDLHLEYAMGKRYFC DYCDRSFQDNLHNRKKHLNGL QHLKAKKVWYDMFRDAAAIL LDEQNKRPCRKFLL\TGQRDFG SN\CRFS\HMSER\DLKEL\SIQVE
						EGRGEPREWLL\DAPELP\EGHL GGLGLEERRAKRLELQPQVSQ G*NPFRTT/VSFQ*PRRVGQPSS GSLPPNSCGHPPPGGLALLQPR V\QWG
19951	50319	Α	20067	2	418	
19952	50320	A	20068		1134	ATTARRRGAPGTPSLAAEGNGL SARVDEGVEEFFSKRLIQQDRL WAPEEDPATEGGATPVPRTLRK KLGTLFAFKKPRSTRGPRTDLE TSPGAAPRTRKTTFGDLLRPPT RPSRGEELGGAEGDTSSPDPAG RSRP\PTQEIARPTR*YCCLPRRR QRWVPDPTSGGPWSGEKQNWL HPLNSGYK*CCRG*ASAEAAGV PKARGSKAKMARSRKLAQMV TLWTVPRRPLPSRSSPAPT/PVS ADPSCRPGPGSQGPESATWKTL GQQLNAELRSRGWGQQDGPGP PSPGQSPSPCRTSPSPDSLGLPED PCLGPRNEERPLRLQRSPVLKR RPKLEAPPSPSLGSGLGTEPLPP QPTEPSSPERSPPSPATDQRGGG
19953	50321	Α	20069	3	206	2.12.331 E.E. 131111 QROOO
19954	50322	A	20070	3	349	
19955	50323	Α	20071	226	610	
19956	50324	В	20072	18	1306	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19957	50325	Α	20073	3	1665	LRPQGREGGDWEGGRSPFSLPY
						GSLRGRVVSTAQFRVPMPSASA
						SRKSQEKPREIMDAAEDYAKER
						YGISSMIQSQEKPDRVLVRVRD
		l				LTIQKADEVVWVRARVHTSRA
		İ				KGKQCFLILRQQQFNVQALVA
						VGDHASKQMVKFAANINKESI
						VDVEGVVRKVNQKIGSCTQQD
						VELHVQ\KIYVISL\AEPRLPLQL
						DDAVRPEAEGEEGRATVNQD
						TRLDNRVIDLRTSTSQAVFRL\Q
						SGICHPLRETLINKGFVEIQTPKI
						ISAASEGGANVFTVSYFKNNA\
						YLAQSPQLYKQMCICADFEKVF
	1					SIGPVFRAEDSNTHRHLTEFVG
						LDIEMAFNYHYHEVMEEIADT
						M\VQIFKGLQER\FQTEFQTVNK
			ľ			QFPCEAISNFWEPTLRLEYCEAL
						AMLRGKLESEMG\DEDDLSTPN
						EK\LLG\HLVKEKYDT\DFYILD
						KYPIGL*RPF/HLYPCP*PPRNPQ
						QVPTLTNMFHEREKKYLSGAQ\
						RIH\DPQL\LTERALH\HGI\DLEK
						IKAYIDSFRFGSPSP/HAGGGIGI/
						ENRVTMLFLGLHNV\RQTSHVP
19958	50326	Α	20074	3	424	
19959	50327	Α	20075	178	859	
19960	50328	С	20076	304	592	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
19961	50329	A	20077	1381	2936	PTNPRPSPGLARWAESYEAKCE RRQEIRESRRCRPNVTTCRQVG
						KTLRIQQREQLQRARLQQFFRR
						RNLELEEKGKAQHPQAREQGP
						SRRPGQVTVLKEPLSCARRISSP REQVGRVKGHTAVCSGTLSGQ
						GGWV/R/RPPSLHRACRES*DNC
						PSSLALTLSHPIHSSLERTQ*I/SH
						ALTFWNSMIYPRSSRLILPQASD
						SSW/KGKGMKFFLNKAWASMC
						SHLIFLSTGKHSEGYFHLRNLEL
						NEKNVFEP\KAKESLQKVLEEK
						MNAEQQLQSTQVWGCHIDMG
						LGTPGSLGNKGSQLQRTLGTQH
						RELESQLHVLQSKLQVPGTGGG
						EGRQGMGRRDGDERSCSAKIE
						CLQGDRDLCSLDTQDLQGTLLL
						GGLECMAAMAK*AKKKETEAF
						QILKVHRVFLGSWPTKGTQVW
						SPKSFPNEVEPEGTGKEKDWDL
						RDQLQKKTLQLQAKEKEVRG*
						PEIVRAHLRNSRETWEAVKNCR
						EELNQSQQLPPRVRGSLLPIKP
100.50		<u> </u>	20052	1	1150	WMMGRFLL*PSFAHKDNLMI
19962	50330	В	20078	1	1173	SSRFLFSLGWLWEAPEVGTRSG
19963	50331	·	20079	16	1584	WLQRRPGLGAG\LGG\QEKQFL
						PALLSFFIYNPRFWPREGEEENK
						ILFYHPNEVEKNEKIRNVGLCE
						AIVQFTRTFSPSKPAKSLHTQKN
						RQFFN\EPEENF\WMVMVVRCPI
						IEKQSKDGKPVIEY\QEEELLDK
						VYSSVLRQCYSMYKLFNGTFPE
		1				SHGRRR\VKLLKERLEKFFHRY
						LQTLHLQSCDLLDIFGGISFFPL
		İ				DKMTYLKIQSFIN\RMEESLNIV
						KYTAFLYNDQLIWSGLEQDDM
						RILYKYLTTSLFPRHI\EPELAGR
						DSPIRAEMPGNLQHYGRFLTGP
						LNLNDPDAKCR\FPKIFVNTDDT
	1					YEELHLI\VYKAMSAAVCFMID
						ASIHPTL\DFCRRTGTASLGPQL
				]		HSGWASGHLVEQF\NINKGDVP
						GLRK\EPQFKFI\YFNHMNLVEK
				1		STVHIEENAPACPLTSVHPDL\M
				1		KIPR*PSNSDLYPRV\DEDEGDH
						C*RALSDYW\VVGK\KSDPAGS
						LYVIFEIQKNAKP*LKVK*KRSK
<u> </u>		<u> </u>		<u>L</u>		KPLCNASSNNILPSLD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	• · · · · · · · · · · · · · · · · · · ·
	sequence	Ì	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19964	50332	Α	20080	85	420	LSNLFLHATNDGALTSTTIPHPP
						MRQPTS/S/GDGSEIFLHTELKLL
						EPPHAWPQSCPPRL*MTRQQLL
						PMTFLLKAHSKAGIESPGLNPR
						PGFSLHPVLPAASSAAFLLPSFR
19965	50333	Α	20081	220	563	
19966	50334	В	20082	1	576	
19967	50335	Α	20083	111	3586	TMFGLKPPLYFLPGSSLSSESSP
}		1				VSSPATNHSSPASTPKRVPMGPI
		ł				IVPPGGHSVPSTPPVVTIAPTKT
l.		ł				VNGVWRSESRQDAGPRSSSGG
						RERLIVEPPLPQEKAGGPAIPSH
						LLSTPYPFGLSPSSVVQDSRFPP
						LNLQRPVHHVVPPSTVTEDYLR
	1	ŀ				SFRPYHTTDDLRMSSLPPLGLD
						PATAAAYYHPSYLAPHPFPHPA
						FRMDDSYCLSALRSPFYPIPTPG
						SLPPLHPSAMHLHLSGVRYPPE
						LSHSSLAA
19968	50336	Α	20084	576	1261	GDDLCSVGAPCRSTWACSTPLR
						CVRRSREFPGLPHAAGAARGFR
						LPAGRPVPARPGAAAARAAAA
		ŀ				GEAHRPLPCGRRPGPVRRAAVP
		ļ				PRPRPPRRGLRGRREGPAAPAG
						PGGPSLLCASAGRHGPPGLRAG
İ						QGLRDSAGEAGGEPAP/AGPRP
		i				EASP*RSSLCENTCTEAGLGPRR
						AAGDLGSPWLTSGLLWGSFFLF
}						FLFFFVILRQSLLLSPKTEVAVA
,						QLMSHCSL
19969	50337	Α	20085	248	356	EGEEEGDEENDPD\MTQRRIKT
						QQSASSSEAGCMWP
19970	50338	Α	20086	1043	1191	NFYFSCKNKNKANKQQITWK\
						KGLSNGCKFILSGGRQKGRTLP
	}					QRERMA
	I	1	I	I	L	1 \

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	, · , ·
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			'	sequence		
19971	50339	Α	20087	287	1401	CQNPQIPCSPFKKRLDG/LWVET
						PT\GYIESLP\RVVKRR\VNALKN
						LQVKCAQIEAKFFE/EKVHGFFE
						RKYAVL\YQPLFD\KRF\EIINAI
		1				YEPTE/EKECEWETR*RKDEDF
						GGNWKEKGQIEDEKKDE\EKE
						DPKG/IFPEFWLTVFK\NVDLLS
						DMVQEH\DEPILKH\LKDIKVKF
						SDA\GQPMSFVLEFHF\EPNE\YF
						TNEVL\TRTYRMRSEP/DTRLIL
						DPFSFDGPEIMGCTGCQIDWKK
						GKNVTLKTY*GGGRKHKGRGT
						\VRTVTKT\VS\NDSFSNFFPP\PE
		İ				IPE/DVGDLDDDAGSLFLAGRLS
						EIWFTFYRERIIPRSSVNILPGEA\
ŀ						IEDDDDDYDEEGEEADEEGEEE
						GDEENDPRLLTQR\RIKTQQSAS
19972	50340	Α	20088	1	350	DNLEAEAVAPCTAAARAVAPD
						LFVFAA*PESVSAGGPQQPCRSP
						IVKPGLPSFRPSSCTQPWLTHSW
						AEDVDLRVNFAMNVGKARGFF
						KKGDVVIVLTGWRPGSGFTNT
		<u> </u>				MRVVPVP
19973	50341	A	20089	1	399	
19974	50342	A	20090	1	394	
19975	50343	В	20091	82	351 789	CGTSRMCHTTEELA/SDPIL/YRP
19976	50344	Α	20092	3	1/89	1
						LAVALDTKEPEI/RNGLIKGSGT SEVELKKGATLKFTRNNAY/ME
						KCDENILWLDYKNICKVVEVG
İ						SKIYVDDGLISLQVKQKGADFL
		İ				VTEVENGGSLGSKKGVNLPGA
						AVDLPAVSEKDIQDLKFGVEQD
			1			VDMVFASFIRKASDVHEVRKV
ł		1				LGEKGKNIKIISKIENHEGVRRF
						DEILEALGHWAVVPLKPTLALA
						LTCFSSSLGLSSLHLSPPSTQLSC
						SKHSTLHLPFSPTTAAPPGLLL
19977	50345	Α	20093	3	424	OLLIGIDATION THRU TOLDE
19978	50346	A	20094	2	173	IYHLQLFEELR/RLAPITSDPTEA
		1				TAVGAVEASFKCCSGAIIVLTK
						SGRSAHQQELGP
19979	50347	В	20095	1	1362	
.,,,,	150577	ــــــــــــــــــــــــــــــــــــــ	1-00/3	1	1-302	<u> </u>

SEO ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
19980	50348	A	20096	1	1982	QRDNLEAEAVAPCTAAARAVA
						PDLLRLCSVRPSRSAPEDRQQP
			:			CLFSHSEAGTAFIQTQQLHAA\
						MADTF/LWEHMCRL\DIDSPPIT
						ARNTG\IIC\TIGPGLPDQVETLK
						EDD*SLGMKCGSVLNFSSWEL
						HGVTMRETIQGMWRTANGKA
						FAFLTPFLLPGPLAVGFRTLKGP
						EIPNWGSSKGNGT\AEVGA*RK
						GATLKFTL\DNAYM\EKC\DENI
						L\WLDYKNICKV\VEVG\SKIYV
						DDGLISLQGKQK\GADF\LVTEV
				}		ENGGSLGI\MKGVNLP\GAAVD
						FACLCRRKDIQDLKFG\VE\QDV
						DMVF\ASFIPQSILMFHEVRKVL
						G\EKGKDHQGLSSKIRESMRGF
						RRFD\EIL\EASDGI/LWVARG*F
						*ALRIPAEK\VF\LAQKMIDLDRF
						NRA\GKP\VICATQMLESMIKKP
						RPTRA\EGSD\VAN\AVLDWKPT
						CIMLSWEKQPKGDYPL\EAV\R
						MQHL\IAREARWLPSYH\LQLF\
						EELRPPGAQLTSDPTEATAVG\A
						VEASFKCC\SGAII\VLTKSGKVC
						SPRLARYRPRAPIIAVTRNSPGK
	1					LVRAHL\YRGIFPG\VCQGPDSR
						RAWA*DIG\LRVKLCHGMFGK
						ARGFFQERGDVVHCA*PEWRP
						WLPAPPNNMRVVPCCRDGPRG
						PSSSRPCSHPFFPKPIPFRPSNGL
						VELHSGAVNVATG

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19981	50349	A	20097	215	1893	TVTFSQCRFGKMLPLEKAFASP QELPSPAGSAHAGVSSRSSSRK NPRPSLRGPLLTLEFSRLRFREF VY\QEAAGPHQTLARLHELCRQ WLMPEARSKEQML/ESLLVLEQ FLGILPDKVRPWVVAQYPESCR SSLLVEGLADVLEEPGPSTVDR DAAGLPCGLILNS*RWSVREAH GPSAATRRAREPQPGPWSWGD PGTF*DTLAFSGPPVSGTEEGQ/ MKQRPKRTALP/HTEQVGGHE QVGGRSQ/RTTNVSEAEVLSRG PSAPGGVGPPGPC\EENLKSYRK LLLWEGGRRQESAGCACEEAA PAG\GCLSCLRRPLGTPLPIPRR A\H*GGGRAAWEGPGPAGPPGR GVRLCHRIAEAVRHPAACP/ET GARRNWEPRGRTPRMGTGRAS RA/CSSSGDSAGLEAGQGPGAD EPGLSRGKPYACGECGEAFAW LSHLMEHHSSHGGRKRYACQG CWKTFHFSLALAEHQKTHG\KE KSYALGGARGPQPSTREAQAG ARAGGPPESVEGRGSPRTPRGA EVSRCAVPFRRGRFAGRESQTR
19982	50350	Α	20098	5	372	HWAFPRPWVAASCGVCGRPLP
19982	50350	A	20098	1	129	
19984	50352	A	20100	<del> </del>	627	
19985	50353	A	20100	1	441	
19986	50354	В	20101	1	477	
19987	50355	A	20102	2007	2156	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
19988	50356	A	20104	1	717	MKLRTLAVSVTALKVARLGSA
1						PSDVQMCLEFLPSDSGAQLASP
						SGSRTGAAGGAACQSGAVRSY
						SSALGWSMGLGAVEQGVVLV
						GDARAAQEPMEWMGGSGMAG
						CRSRALHRGKAAKARREIEHSA
				1		GRKGLFGSARLIPATAMAPRSR
						LLSLGRRGNFRSRVLRRKSRPL
						EEAARRWRDCPTGFGALVAGA
						GSGRAPGVPPKRLPARTKAQVS
						RETRGSGEVTAQGGKGPRGLW
						ILLWGFYNLLPTTSHAESYPERL
						IQKVVSEKIARALARYLHLPFG
						l -
						LLGGNKTIVYSWEALRFHHQLT
						PSVITNTVIKVYEEPKLSQQKSR
						TLDVSTDEEDKIHHSSESKDDQ
						GLSSDSSSSLGEKELSSTVKIPD
				1		AAFIQAARRKRELARAQDDYIS
				]		LDVQHTSSISGMKRESEDDPEIS
				1		RNEETSEESQEDEKQDTWEQQ
				İ		QMRKAVKIIEERDIDLSCGNGS
				1		SKVKKFDTSISFPPVNLEIIKKQL
						NTRLTLLQETHRSHLREYEKYV
						QDVKSSKSTIQNLESSSNQALN
						CKFYKSMKIYVENLIDCLNEKII
						NIQEIESSMHALLLKQAMTFMK
						RRQDELKHESTYLQQLSRKDET
						STSGNFSVDEKTQWILEEIESRR
						TKRRQARVLSGNCNHQEGTSS
						DDELPSAEMIDFQKSQGDILQK
						QKKVFEEVQDDFCNIQNILLKF
						QQWREKFPDSYYEAFISLC*RQ
19989	50357	Α	20105	154	744	SQRNLCLGRVWYILFFLFAFLQ
						TLELVSHERRGRTGITNA*VSNS
						PRRELGFHVL*R*QFLLLFLFSH
						RLRSPAGFT*WIPHRGCRWSCL
						PVLRHALAFLSPWVVDGTGRR
						GAGGGARRGGSGHTGAHGVG
						GRLRHGRLQVPSPLRARSPFLLS
						CTLSCQLDSPCLTCSQGHQYKY
						QHSDKGKRLTIVTKSFDFIPS
19990	50358	В	20106	1	531	
19991	50359	С	20107	99	323	
19992	50360	Α	20108	305	658	
19993	50361	В	20109	79	936	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
19994	50362	A	20110	381	1169	LPSRGAGLGTCSPPCLSLPPTPW APVRPEPPQGAPPPAPRRPVPST TQGLRSASARHGTGRQLHLQP WCGIHWVKLAGLLSLVGSQHL
						RGQGSRAAAALTHMHS/PHAC THAHTRAEPLRLPPPPRPLPPPL
						SPPPLRLCCLRSLFPCPLTESAA
						KAGSKKFHSVQEIPSPAVCELG
Ì						VGTVLHIDPEKWRAGRENVET
	}					GGKRTGSGKDSTGRGWGQTD
						YGLGKKERTEEASRSCICLVCV
						SMRRDALCGQSVGSHFCQWSL
19995	50363	A	20111	3	4804	KRLENIQKTLEVAFSEAVWMQ
						PSVVLLDDLDLIAGLPAVPEHE
						HSPDAVQSQRLAHALNDMIKE
						FISMGSLVALIATSQSQQSLHPL
			:			LVSAQGVHIFQCVQHIQPPNQE
						QRCEILCNVIKNKLDCDINKFT
				}		DLDLQHVAKETGGFVARDFTV
				ŀ		LVDRAIHSRLSRQSISTREKLVL
		1				TTLDFQKALRGFLPASLRSVNL
						HKPRDLGWDKIGGLHEVRQIL
						MDTIQLPAKYPELFANLPIRQRT
						GILLYGPPGTGKTL
19996	50364	Α	20112	2	1844	
19997	50365	Α	20113	746	1215	PNINRFLKHYREPVPKKRERNY
						STLWPLCPSTLNKIYTIMSYLSP
						ATKEYPFFLLLLEQEC*AE*LLA
						LAVSQPLLSSTTNCLKN*MVTW
						NGSLIPWSPCKINLTP*/PAVVL
						QNRRALDLLTAESGGTFLFLEE
		<u> </u>				KCCCYVNQSGIITEKVKEIQGRI
19998	50366	A	20114	1	1338	
19999	50367	В	20115	1	4935	
20000	50368	Α	20116	3	421	
20001	50369	С	20117	33	571	
20002	50370	Α	20118	3	323	
20003	50371	Α	20119	146	809	EKCRHNCSSRVWQSLVSQSVW
						ATEGQYGRTKNARPVQVK\DS
						ASFPYQRRYPLRLEAQQGLQKI
						VKDLKAQGLVKPFNSPCNTPIL
						GVQKPNGQWKLVQDLRIINEAI
						VPLYPAVPNPYTLLSQIPEEAE
						WFTVLDLKDAFFCIPVHRESQF
						LFAFEDPSNPTSQLTWTVLPQG
						FRNSPHLFGQALAQDLSQFSYL
						NTLVLRYLDDLLLAAHLETLCH
20004	50372	A	20120	3	1147	
20005	50373	Α	20121	2	1658	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20006	50374	A	20122	2	211	MGIYDFSYFPPGCCLKSRRKSS
	İ	-				RGSQE*LALSPGFSGSFGSIQDL
						TQV*CIQDSTPATLTAVGVDKM
20007	50375	В	20123	86	586	
20008	50376	A	20124	66	433	SSSRTEGARGKRQPMPSP/STEP
						RVCLTIEGQEVNCLLDTGAAFS
					:	VLLSCPGQLSSRSVTI*GVLGQP
						VTRYFPQPLSCDWGTLLFSNAF
						LIMPESPTPLLGRGILAKAGAIIH
						HAVLVTAAL
20009	50377	A	20125	1	459	MAVRYTDENVLRKGTREAGT
		1				MMRLRGTTECCADFAAWDVT
						HDALRATGGELPGAQVMLTTT
ŀ						ECGRHVDFCDRAVWLPRMWG
						YPLALPGEMRKLYTVRMAGRD
						ILAKAGAIIHLNIGEGTPVRCPL
						L/EGINPEVWKTEGQYRQAKNA
						RPVQVKL
20010	50378	A	20126	260	392	VEVVILPMPVPNLPSIPARITKG
	30370	'	20120			RMGTLLLRGLGTT*LYKSCSV
20011	50379	A	20127	1	1740	
20012	50380	A	20128	1	858	MEWEDNLPLELGRTVAKLLSD
						HSQTPLGIQMFLLFSLSLRKSPL
						VCLSYLFNFRFTLESEVQHLSG
		1				AITLTAWPKIPFLGIREAKSPRS
						ENTRLATILEAGHRHLGTSVSK
ľ			ļ			DHPVTFWRPRRDLQSDLKQIKI
ì						DLGKFSDNPDGYIDVLQELGQS
						FDLTWRDIMLLLNQTLTPNERS
						ATITAAREFGDLWYISQVVAAV
				İ		AGLVSEAVKIIQG\LTVWT/SHD
						VNGILTAKGDLWLSDNHLLKY
						QALLLEGPVLRLRTCATLNPAT
						FLPDNEEKIEHNCQQVIAQTYA
20013	50381	A	20129	1	6530	MWGSDRLAGAGGGGAAVTVA
ŀ						FTNARDCFLHLPRRLVAQLHLL
						QNQAIEVVWSHQPAFLSWVEG
						RHFSDQGENVAEINRQVGQKL
						GLSNGGQELHAVSLEQHLLDQI
						RIVFPKAIFPVWVDQQTYIFIQI
				1		VALIPAASYGRLETDTKLLIQPK
	1					TRRAKENTFSKADAEYKKLHS
		1				YGRDQKGMMKELQTKQLQSN
						TVGITESNENESEIPVDSSSVAS
[						LWTMIGSIFSFQSEKKQETSWG
						LTEINAFKNMQSKVVPL
20014	50382	В	20130	1231	2065	
20015	50383	A	20131	68	372	
20016	50384	A	20132	1	1461	
		<u> </u>		L	L	L,

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	Nucleotide location of first	codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
20017	50385	A	20133	1	806	AVWCTHCCAPTPSGPVLPHSAA MSFLKSFPPPGPAEGLLRQQPD
						TEAVLNGKGLGTGTLYIAESRL
						SWLDGSGLGFSLEYPTISLHALS
			1			RDRSDCLGEHLYVMVNAKFEE
						ESKEPVADEEEEDSDDDVEPITE
						FRFVPSDKSAF*FPCSVEAMFTA
·						MCECQALHPDPEDEDSDDYDG
						EEYDVEAHEQGQGDIPTFYTYE
						EGLSHLTAEG\QATLERLEGML
						SQ\SVSS\QYNMAG\VRT\EDSIR
						DY\EDGDGRLDTTP\TVCLGQF\
						EDADV
20018	50386	Α	20134	225	420	RLLGDVCGRLRPRGHEEQRPG
			:			GPGARSHSRSGPRMGRDCAGA
						VPRVTPK*SACAAGGRGGPAT
20019	50387	Α	20135	1	687	
20020	50388	Α	20136	1	2979	MLRRKKQEDWAAKGSSHACG
						QIPEEREKGFAVGNVVANLGLD
		ľ				LGSLSARRFRVVSGASRRFFEV
						NRETGEMFVNDRLDREELCGT
		1				LPSCTVTLELVVENPLELFSVEV
		l				VIQDINDNNPAFPTQEMKLEISE
						AVAPGTRFPLESAHDPDVGSNS LQTYELSRNEYFALRVQTREDS
						TKYAELVLERALDREREPSLQL
		1				VLTALDGGTPALSASLPIHIKVL
		1				DANDNAPVFNQSLYRARVLED
		1				APSGTRVVQVLATD
20021	50389	A	20137	2	296	
20022	50390	Α	20138	1187	1292	
20023	50391	Α	20139	2	3250	LTSRWRPWPRPPTSASSAGWFC
		1				ASTGPWTAAPAKAPPSWASWT
		ł				SRALRSSRPQTSRLWRR*PRSR
						AATPSSSGRGTCGIRPTSVFSTT
						RARSTTRPTSG**KTWTL*MTT
		1				SQ/RLLHQSTDRLTAEIWKDVE
						GIVGLEQVSS\RATAHQVAAPV
		İ				GVCSGQWDSS/HKESLSRLMAT
		İ				LSNTNPSFVRCIVPNHEKRAGK
						LEPRLVLDQLRCNGVLEGIRICR
		ļ				QGFPNRILFQEFRQRYEILTPNA
20024	50392	В	20140	1	1816	IPKGFMDGKQA
20024	50393	A	20140	500	866	QVCLYPGWQHSPGRQRVSG*A
		'				HLHTAPLWICQ*SGDCDWHES*
						GGGM*CVR\FEPPRTSSPPSATS
						GSLLK*KEKTGLSNSPCSTSTSS
1		1				GGRTPSTDRMGKTRRQYPSGA
						GNSEDKVGFPQVF
20026	50394	Α	20142	130	226	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20027	50395	Α	20143	846	1555	GRKTPSQGRRWSSPRVFLLPGG
20027	30373		20112			PKPCYTYPLVGEKLLFRVLPLFP
						HKPSASNLEKVSHCPECGKTFS
						RSSYLVRHQRIHTGEKPSQVPV
						ECGKGF**ALPT/CTAH/HYGTH
						TGERPYQCGQCGKSFNQSSSLI
·						VHQRTHTGEKPYQCIVCGKRF
						NI\SSQFSAHRRIHTGESPYKCA
			•			VCGKIFNNSSHFSAHRKTHTGE
						KPYRCSHCERGFTKNSALTRHQ
		İ				TVHMKAVLSSQEGRDAL
20028	50396	A	20144	93	764	AQERKALKSTPSHSSFSR/WLSR
20028	30370	Λ.	20177		704	IILKNGSSSSSSSSSSSSSLRE
						RERKSCENVNITRNRR/LTAETI
						ADTIDTSNGSVFLIWTER*EWN\
}						WLPKPWHQYQLQARASLSEEIL
						NNRWEQDPEAFRQDPEDQARS/
						QWLPRGGSAPV\KQRLLLHHDS
						VPDHAFLRTRAAGQDLPWETV
						RRPPDNPDLAPADFSLFPCLGK
						LLAERRCFSLQRRVPWPPENQP
						PHRO
20029	50397	A	20145	1	433	TINQ
20029	50398	C	20146	312	443	
20031	50399	A	20147	1	94	
20032	50400	A	20148	2	122	DLLCPQMG*GWKLTALSLQCS
20032	30 100	, · ·	20110	_		LQDGIERSRAKASQCCLSI
20033	50401	С	20149	25	328	
20034	50402	C	20150	157	309	
20035	50403	В	20151	33	2996	
20036	50404	c	20152	167	274	
20037	50405	Ā	20153	3	130	AGAVQFQALGVPNKDTKSLEL
						HQKPKPR*GQDDNHRMSPVYV
20038	50406	A	20154	2	453	
20039	50407	Α	20155	1	1317	
20040	50408	Α	20156	1	1350	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		l		sequence		
20041	50409	A	20157	970	2559	FPLAYSLLFPP/CSRLSRELLEVV
						EPEVLQDSLDRCYSTPSSCLEQP
						DSCQPYGSSFYALEEKHVGFSL
						DVGEIEKKGKGKKRRGRRSTK
						KRRRGRKEGEEDQNPPCPRLS
		1				RELLAEKEPEVLQDSLDRWYST
						PSVYLGLTDPCQPCRSAFYVLE
						QQRVGLAVDMDEIEKYQEVEE
						DQDPSCPR*L*AITDA*FCVDT
						WRCQVQGKQECVQFHVFNEG*
						ITPTDIAVGFHCSRCLGFHFFLP
						LSFTHLL*VDHTSKAVWQLHGI
						LSKFMENY*AHSFHDHCSLCVP
			İ			RALTQSVL*PLHQCVTRPIR*AH
						FLLSLSLPLPVFLFHSFLPGPGLS
						QHKGNNSLPH*WICPFSF*TVPY
						VSHEI*LGLCGF*FPLAYSLLFPT
						/CSRLSRELLDEKEPEVLQDSLD
						RCYSTPSGYLELPDLGQPYSSA
1						VYSLEEQYLGLALDVDRIKKD
					İ	QEEEEDQGPPCPRLSRELLEVV
						EPEVLQDSLDRCYSTPSSCLEQP
						DSCQPYGSSFYALEEKHVGFSL
						DVGEIEKKGKGKKRRGRR
20042	50410	A	20158	2790	5320	LHLFCHKRQHGGISRALGSSKK
20042	30410	1	20130	2,750	3320	PEMNILEISETLRPQLAEKKQQF
					}	RSLKEKCFLTQLAGFLANRQKK
					}	YKYEECKDLIKFMLRNERQFKE
						EKLAEQLKQAEELRQYKVVVH
						SOERELNOLREKLREGRDASRS
						LYEHLQALLTPDEPDKSQGQDL
						QEQLAEGCRLAQHLVQKLSPE
		İ				NDEDEDEDVQVEEAEKVLESS
						APREVQKAEESKVPEDSLEECA
						ITCSNSHGPCDSNQPHKNIKITF
		l				EEDEVNSTLVVDRE
20043	50411	A	20159	394	3251	GRRVCLFSEDEECKDLIKFMLR
						NERQFKEEKLAEQLKQAEELR
ľ		ľ				QYKVLVHAQERELTQLREKLR
]		ŀ	İ	-		EGRDASR\SLNEHLQALLTPDEP
						DKSQGQDLQEQLAEGCRLTQH
						LVQKLSPENDNDDDEGVS/QVE
						VAEKV\QKSSAPREMQKAEEKE
						VPEDSLEECAITCSNSHGPYDSN
						QPHKKTKITFEEDKVDSTLIGSS
						SHVEWEDAVHIIPENESDDEEE
						EEKGPVSPRNLQESEEEEV\PQT
						PWHE*LSHLSIL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide sequence		in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20044	50412	ΙΔ.	20160	245	641	  ILSVLIAVLCVVGKIPLKEPLCP
20044	50412	A	20100	243	041	
						AAGSAAEACVGQGMPVIQVPC
						SQQRPSSMCWPRNACQSSRAL
						GGDESYVPKNRKSFTKHPVSSC
ļ						/PSPSLGWGRG*R/PGGPGRSPG
	70.440	<del>  </del>	00161		104	LALGFPSFSGPAGFGQGTSSPRF
20045	50413	A	20161	7	194	RPLVESRGIPRSPAASFSTACPR
						NTASIPSAPSWWETAWTQTSS*
70016		<u> </u>	20160	222	500	APPVA*RPS*PSPESPL
20046	50414	Α	20162	232	509	MGVLMHRRCTKAPAWAVATY
						EAPLHCQQLLGVVGVLAGDPS
						LLDLGGIHQQPQGLPITVLHQA
						EEADLVLLRAVTDAAALPVEQ
	1	1				HRP*GAPGLLPPPDGGQ*SGDP
						AAAGGYLQGPRGLGPQPGHPP
						HPGAAGSAGEPRKWLPPRRGL
İ						WCSACA*GPPSTGPGEQRGPGD
						SG*FISASTSASRATSACRSAFIT
						SQTEPPTWPSLARPQSSGLARPP
						APGRTAAGRD**TLPPGK
20047	50415	Α	20163	3	859	SLGGNKSRPTRRCCSGSRCRCR
						DKSAAVSGALWRGCRESPRAP
						PPAPQTIPRAWPSPTAVLTHPLL
						AGPSRWGSSCTGAAPKPPPGRG
		١,				PLRGSPALPAAPGCGGCPGWGP
						KPLGPWRYPPAAAGSPDHCPPS
						GGGSRPGAPQGGNRCSRSARRT
						APPSCGCSEVGSL*WSPRSGPRP
						PGWFGWAGACGGSGGSFAASA
						ALAWEGPEQSQEALQPQSPQRP
j						APAAAAAGAGPRLSSPRRRAW
						SSPG*/HGIPWPTHASAALPAAG
l						HRGSFSGILP*RAGKRSAMLST
20048	50416	В	20164	179	438	
20049	50417	A	20165	1	930	MESPWRQWRLRKVARSGGGR
ŀ						LSNKRKLHLKKKKKGERGGGG
						AEREKAAAAVAPGCKAGAGN
						AKKFVQNGPGTKESSASQVPTG
						PSAAGLLEFARGPLQTLFAWVS
						AAVSAEPRIFMIRECCCLIVPLE
						VLSQRSTWPCEFLISDCCASNQ
	1					QDSMGVGPSEPGAGYNLLVRH
						FLSPSEKCSIRVGVARFSRVLPP
	}	1		1		PLLGISAARAAGGAARQAAQL
		1		1		REGSRCPRLTDIPHAPSTPPEQG
	1					i -
	1					HSAEGAAKEEPKRRSARLSAKP
			1		1	APAKVEAK/PKKAAAKDKSSD
						KQVQTKGKRGAKGKQAEVAN
	1.0.410	ļ	20155	1.00	410	QETKEHLPAV
20050	50418	Α	20166	199	419	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
20051	50419	В	20167	969	1313	
20052	50420	C	20168	481	738	
20053	50421	В	20169	56	3476	
20054	50422	A	20170	1	2679	
20055	50423	A	20171	1	1467	
20056	50424	A	20172	208	379	NPCKKGVLIFQF*LKIHKRQFIC
2000						ADLEFVISE*CTRYPIVAFNWCK PSLLRRSTPVK
20057	50425	A	20173	1	921	I SEEKKS II VI
20058	50426	A	20174	197	355	
20059	50427	A	20175	1	3180	
20060	50428	A	20176	1	1309	
20061	50428	A	20177	1	2295	MFYNDVKKQQQMAALTKEKD
20001	30429	A	20177	[	2273	QLSQSIVMYKSLFEASQQLLNE
]						MKCQVEEARLKEAQLRNELKI
į						HNIDIPTTQQVPHIEALLKRKLS
						EQEELKKKPRRSCTLPNYTKGS
			1			_ ~
						GDVLGKIAHLAQIEDDRAAMVI SWHLASDMDCVVTLTTDAARR
						IYDETQGRQQVLPLDSIYKKTL
						PDWKRSLPHFRNGKLYFKPIGD
		į				PVFARDLLTFPDNVEHCETVFG
		Ì				MLLGDTIILDNLDAANHYRKEV
	<u> </u>	<u> </u>				VKITHCPTLLTRDGDR
20062	50430	Α	20178	3	2550	
20063	50431	Α	20179	256	3680	NRGLLFCFLFLHLHLKEHERHL
						ATVYVFPILSCTIYRSVLLTSSQ
						TIIYTHVGDREAQAALKLGRWS
						HPRTPNAVGAPGPPEGAGGGD
						AVTSQSALLTFSRTRFASGAHA
						GAHPVLLRNEEEKGAPALVAPI
						FSAEGPTCSLWWTLRPASTAGL
		ł				KLPAAASTRPSQRGLTADGGGP
						GGASVGTEEDGGGVGHRTVYL
						FDRREKESELGDRPLQVGERSD
						YAGFRAYETVKDGVTLYLLQS
						VNQLLLTATKERIDF
20064	50432	Α	20180	8	267	RGEMPGERLKAEVFLPPSPLSPT
						QVVGPLHRVWS*IGNASSQRGE
						AQVRS*VSFSLRLEDSTPWIPGT
						RALRGCWLWRDFQTLQSG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence	:	
20065	50433	Α	20181	1	3692	MNPIREMQVSGIALKGTMKGW
20003	30433	( )	20101		3032	TELPKSPVWTQGNKGRNYFTN
						LALSLLFEDDTKLNPYAGGDAI
				1		RVRISVGPFKGERFTVPGLASEC
						GKNSFLRFPSNSRRKQLCGRVH
						GSFLWFSMMSQSMGGDNLSSL
			Ė			DTNEAEIEPENMREKFFRSLAR
						LLENKSNNTKIFSKAKYCQLIK
						EVKEAKAKAKKESVDYRRLAR
]		1				
						FDVILVQGNEKLIEAVNGETDK
		1				IRYYLHSEDLFDILHNTHLSIGH
		<u> </u>	22122		1.01	GGRTRMEKELQAKYK
20066	50434	Α	20182	24	131	PVHKRWS*NDECYRHCTV***Q
		<u> </u>				PGSEENNPALENR
20067	50435	С	20183	99	281	
20068	50436	Α	20184	1	4200	MVILKDFMKFEMNSEEGDFLR
		ļ		}		TCDFFFLRHLVAVKQLAHVKV
	1					DRFFPRRQSLAESRTAGIRLQTN
	1	1				VQKMVSSFTTSDRKQTSPSSYS
						VSFIPEFHFLGPPTHCPQTAVSQ
		1				CDLLFQLHSPAEALFPARTLPG
		l				GPRSPHLRVPAAPAPLQHDPRH
						TGPPLPPTANRGATPPLPLGPGA
						LERSGRLGELNSASSFRLRRDN
						PQSPAARPRGSHGLAGIEGERT
						LINYYCQERYFHHVLLVASEGI
						KRYGSDPVFRFY
20069	50437	Α	20185	1	1464	
20070	50438	Α	20186	335	855	RIASMPRLPPRTQGRGSARLPSR
		1				ARPGLHWHHRHTMLPPKPRRW
						R/RALGPAEAGMLEKLEFEDEG
						ECAPPAAGPALPRPGRPRTGTR
						GDPAPHLRRCPTASCVLRKPRG
						RGGNRGGDAGAGELGLRGPCD
						LPSQNFLCFLVDVLKNGLLALQ
						LFSSNPTASCAESLPMLRTPVI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			-	sequence		
20071	50439	I A	20187	2	1160	SARPAASPPAGERASQLMLIAV
20071	130437	^	20107	<u> </u>	1100	LLLPLRRRWRRALGPAEAGML
						EKLEFEDEAVEDSESGVYMRF
						MRSHKCYDIVPTSSKLVVFDTT
						LQVKKAFFALVANGVRAAPLW
						\ESKKQSFVGMLTITDFINILHR
						YYKSPMVQI/YMELEDHKNLET
						\WRELYLQ\ETFKPLVNISP\D\AS
						LFGWLYTSLIKNKIHRLPVIDPIS
						GNALYILTHKRILKFLQLFMSD
						-
						MPKPAF\MKQNLDELGIGTYHN
						IAFIHPDTPIIKALNIFV/EKDRIS
						ALP/VLWDESGKSW*IFNSPNLN
						VN*ILACLEENIPNNLR*STVT\Q
						A\LQHRSQYFEGVVKCQLSWKF
						LETHRWTRIVRAEGPSGLVVV\
				ĺ		NEADSIVGIISLSDIL\QALIL\TP
20072	50440	<u> </u>	20100	50	205	AGAKQKETETE
20072	50440	A	20188	347	205	NDCDCAVANECONDAITCHIOTE
20073	50441	A	20189	347	1581	NPSPC/WAVFFQKDAITSKIQTF LSMLHLYN**/SSPSKENTRTVE
						KLYKCSVDIRKIRRFKKDGVLL
						RR*KPYVERNCRIFYQELGAR*
}						D\AVASI\LERCPEAIV\*SPTAVN
						TQRKLWQLVCKNEEELIKLIEQ
						FPESFFTIKDQENQKLNVQFFQ
						DVGTKKCGH\CRLLTAAPNCFS\
						NPVEKNKQMV\EFFQESYLD\L
						GGSEANMKSFGLLKLFKPKPFY
						FV*ISPTAIKETLEFLQEPRFPPA
						LKFLQLPIPNSKGFLFQLCPKKY
:						TE*YFLL*KCLLNAQIFMDP/VS
				:		N*VLKWPCPFILFLFPVLEERM
						QGLLREGISIAQIR\ETPMVLELT
						PQIVQYRIRKLNSSGYRIKDGHL
						ANLNGSKKEFEANFGKIQAKK
						ALQNLLLRSLNYYDELTVGEC
20071	50440	<del>  -</del>	20100		455	GNRKGRGTNVNTTL
20074	50442	A	20190	3	455	
20075 20076	50443 50444	C A	20191	72 3	218 452	LGILLVDDFVNVLSSLAVYAED
20076	30444	A	20192	]2	432	SEPESDGEAGIEAVGSAAEEKG
		1				GLVSDAYGEDDFSRLGGDEDG
		1				YEEEEDENSRQSEDDDSETEKP
				}		EADDPKE\NSAQ*QRHPGLSKL
						MAQLVGLRGQKNHILNMIRDV
20077	50445	<del>  -</del>	20102	ļ, —	224	TQQPQLSLAAHTPHTPFHLQ
20077	50445	A	20193	1	334	
20078	50446	Α	20194	]	687	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20079	50447	A	20195	1	1280	RPGCGNKMAGKKNVLSSLAAY
	1					A\*ISEPESDGEAGIEAVGSAAD
	}					EKGGLVSDAYGEDDFSRLGGD
	1					EDGYEEEEDENSRQSEDDDSET
						EKPEADDPKDNTEAEKRDPQEL
						VASFSERVRNMSPDEIKIPPEPP
		ļ				G\RCSNHLQDKIQKLYERKIKE
		ŀ				GMDMNYIIQRKK\EFRNPSIYRE
						/RLIQFC\AIDELGTNYPKDMFD
		İ				-
						PHGWSEDSYYEALAKAQKIEM
						GQIGKRPKKERNQKFEFCDGAP
						KKGT/TRTNATSTTTTTCQAQL
	ļ				FADCFRRGKRQVGIRAIPSATKP	
	ļ				KRTPPPKPPPNTLTTTATLPS/V	
	1				VVTVTHPSASGSKDPPVISCCG	
	1					APIVKERPKQVTLRGPPLGLER
1						DRAAPVTHCPVGGAHFVYFRT
		1				GTYSPRCHLRGASVWHSRWKD
		i				RQHGSQALWTGSVHAPPGVCR
	1	1				LLKVPYSYLLASQKK
20080	50448	Α	20196	1	542	MAYSAPEKSCTEGHESPH/HHT
	1	1				AGHSGWHGG*ALRQRGHSCPK
		l				EGTRGHPQTLAP/HHQCPQHAA
		1				GLPR/HNQLLHD/WFRHHHHRH
						WHHHHHHHQ*FPECQPSTSHQ
		l				ANPHHRPDLHDRQAPDSGYEG
						VSPKRVAEANFSLGSAPQAAQ
						GRSHGQSPGVHGFFSHGSPAAS
						SDIMKRLPPPGLF
20081	50449	Α	20197	45	171	RKTRERIGHDVICNLL*C**SS/C
20001	30445	^	20177	143	1171	GCGLS/CICS*IRHSKSSRKA
20082	50450	A	20198	277	467	GLGPHDYLYSILSIERSCCC*CC
20062	30430	^	20196	277	407	CCCCRRRCCCCC/CV*GCSRFL
20083	50451	A	20199	1	1080	CSIAESTPSGALRRLRGGR
20084	50452	A	20200	34	119	
20085	50453	A	20201	1798	1894	
20086	50454	A	20202	363	778	GLGPHDYLYSILSIERSCCC*CC
20000	30434	<b> </b>	20202	303	1770	CCCCRRRCCCCC/CV*GCSRFL
						CSIAESTPSGALRRLRGGRSGGS
				[		GGCCSPGGCREGGRGNTKPAT
		İ		1		
						RCEVICGGGSGGGGARPAAPPG
20087	50455	A	20203	1	360	GPDASRVCWFSLNTSIVFFSCPF
20088	50456	В	20204	1	3681	
20089	50457	A	20205	1	293	
20099	50457	В	20205	38	282	
20090	120420	ր	20200	120	1202	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ĺ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			-	sequence		
20091	50459	Α	20207	1	1124	VDTVKRLLIKKLPPVLAIQLKRF
						DYDWERECAIKFNDYFEFPREL
						DMEPYTVAGVAKLEGDNVNPE
						SQLIQQSEQSESETAGSTKYRLV
						GVLVHSGQASGGHYYSYIIQRN
						GGDGERNRWYKFDDGDVTEC
						KMDDDEEMKNQCFGGEYMGE
						VF\DHMMKRMSYRRQKRWWN
į						AYILFYERMDTIDQDDELIRYIS
ł						ELAITTRPHQIIMPSAIERSVRKQ
		ĺ				NVQFMHNRMQYSMEYFQFMK
		Ì				KLLTCNGVYLNPPPGVAEKTQL
						LKLSVPATFMLVSLDEGPGPPIK
						YQYAELGKLYSVVSQLIRCCNV
						SSRMQSSINDAETAFNKIQHRF
						MLKILNKQGIEETYIKITRAIYD
						KPTANIILNGQMLEALPLRNRT
20092	50460	A	20208	1	1053	
20093	50461	Α	20209	1	7785	MTATTRGSPVGGNDNQGQAPD
		i				GQSQPPLQQNQTSSPDSSNENS
						PATPPDEQGQGDAPPQLEDEEP
ł						AFPHTDLAKLDDMINRPRWVV
		l				PVLPKGELEVLLEAAIDLSKKG
						LDVKSEACQRFFRDGLTISFTKI
						LTDEAVSGWKFEIHVSLAFDFK
						GSDLHSDACEWEERCIINNTHR
						LVELWCGPSCPQDWF\PLLELL
						AMALNPHCKFHIYNGTRPCESV SSSVQLPEDELFARSPDPRSPKG
						WLVDLLNKFGTLN
20094	50462	A	20210	1201	9141	ADKIAKICPVSSMTATTRGSPV
20094	30402	^ <u>`</u>	20210	1201	7141	GGNDNQGQAPDGQSQPPLQQN
						QTSSPDSSNENSPATPPDEQGQ
						GDAPPQLEDEEPAFPHTDLAKL
						DDMINRPRWVVPVLPKGELEV
						KLEANIDLSKKGLDVKSEACQR
						FFRDGLTISFTKILTDEAVSGWK
						FEIHRCLVELCVAKLSQDWFPL
						LELLAMALNPHCKFHIYNGTRP
						CESVSSSVQLPEDELFARSPDPR
						SPKGWLVDLLNKFGTLNGFQIL
						HDRFINGSALNVQ
20095	50463	A	20211	2	710	HEIGHIOSALITY
20093	20403	$\Gamma$	20211		/10	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20096	50464	A	20212		1703	RPLDVGGPAAGTPGVLSRPCPS TAALAPKPFCAAPRPQPDAPAC AGATGGSCADFDSGVDFVRRR SSGLWGPQPPLFPVKNYTEDPV AFKDVAVNFTQEEWALLDISQ KNLYREVMLETFWNLTSIGKK WKDQNIEYEYQNPRRNFRSVT EEKVNEIKEDSHCGETFTPVPD DRLNFQKKKASPEVKSCDSFVC EVGLGNSSSNMNIRGDTGHKA CECQEYGPKPWKSQQPKKAFR YHPSLRTQERDHTGKKPYACK ECGKNIIYHSSIQRHMVVHSGD GPYKCKFCGKAFHCLSLYLIHE RTHTGEKPYECKQCGKSFSYSA THRIHERTHIGEKPYECQECGK AFHSPRSCHRHERSHMGEKAY QCKECGKAFMCPRYVRRHERT HSRKKLYECKQCGKALSSLTSF QTHIR\MHSGERPYECKTCGKG FYSAKSFQRHEKTHSGEKPYKC KQCGKAFTRSGSFRYHERTHTG EKPYECKQCGKAFR SASQLRIHRRIHTGEKPYECKK
20097	50465	Δ.	20213	3	2413	CGKAFRYVQNFRFHERTQTHK
2009/	30403	Α	20213	ا ا	12413	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20098	50466	A	20214	1	2124	MGKNTPMIQLSPPGPFHNSEYR
20096	150400	^	20214		2124	FPVLSVSAFLLYTCEMFQDPVA
						FEDVAVIFTQEEWALLDISQRK
						LYREVMLETFRNLTSLGKRWK
						DQNIEYEHQNPRRNFRSLIEEK
						VNEIKDDSHCGETFTPVPDDRL
						NFQEKKASPEVKSCESFVCGEV
						1 '
						GLGNSSFNMNIRGDIGHKAYEY
						QEYGPKPCKCQQPKKAFRYRPS
ŧ						FRTQERDHTGEKPNACKVCGK
						TFISHSSVRRHMVMHSGDGPY
						KCKFCGKAFHCLRLYLIHERIH
						TGEKPCECKQCGKSFSYSATHR
						IHKRTHTGEKPYEYQECGKAFH
						SPRSYRRHERIHMGEKAYQCKE
						CGKAFTCPRYVRIHERTHSRKN
						LYECKQCGKALSSLTSFQTHVR
						LHSGERPYECKICGKDFCSVNS
						FQRHEKIHSGEKPYKCKQCGK
						ALPHSSSLRYHVRTHTGEKPYE
						CKQCGKATHNGEKPYACKECG
						KPFGSAQNLRIHERTQTHIMHS
						VERPYKCKICGRGFYSAKSFQI
			}			HEKSYTGEKPYECKQCGKAFV
			1			SFTSFRYHERTHTGENPYECKQ
				•		FGKAFRSVKNLRFHKRTHTGE
						K/PL*IHEKTHTGRKHYECKQC
						GKAFTSSSSFQYHERTH/S/GEK
						PYQCKQCAKAFISSTSFQYHER
						THMGEKPYECMPSGKAFISS\SS
						LQYHERTHTGEKPYEYKQCGK
						AFRSASHLQMHGRTHTGEKPY
20099	50467	A	20215	23	265	
20100	50468	Α	20216	40	470	
20101	50469	Α	20217	235	392	

	SEQ ID NO:	Met	SEQ ID NO:		Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20102	50470	Ā	20218	7	1439	EGADKMATSVGHRCLGLLHGV
20102	30170	` `	20210	ľ	1100	APWRSSLHPCEMTALSQSLQPL
		İ				RKLPFRAFRTDARKIHTAPART
						MFLLRPLPILLVTGGGYAGYRQ
		l				YEKYRERELEKLGLEIPPKLAW
				•		SLGSVALYKSVPTRLL\SRAWG
						RLNQVGAAHTGLRRPRFTSL\YI
						WTFGVNMKEARVEDLHHYRN
						LSEFFRRKLKPQARPVCGLHS/V
						GEA*PFPPAGNRTFSCLPSTAPL
						VSSVSGRGRMTRGGGGCLLGG
						GDPALPGSKPLLPFQISPSDGRIL
						NFGQVKNCEVEQVKGVTYSLE
		ŀ				SFLGPRMCTEDLPFPPAASCDSF
						KNQLVTREGNELYHCVIYLAPG
						DYHCFHSPTDWTVSHRRHFPGS
						LMSVNPGMARWIKELFCHNER
						VVLTGDWKHGFFSLTAVGAT\
						NWGSTRIYFDRDRNTNNPRHSK
						GSYNDFSFVTHTNREGVPMRK
						GEHLGEFNLGSTIVLIFEAPKDF
						NFQLKTGQKIRFGEALGSL
20103	50471	Α	20219	170	1144	ARTRRLAPSPGCRRGPAVKKM
						RHEELLTKTFQGPAVVCGTPTS
						HVYMFKNGSGDSGDSSEEESH
						RVVLRPRVKERHKSGVHQPPQ
						AGAGDVVLLQRELAQEDSLNK
						LALQYGCKHSE*LTMA/NVADI
						KKVNNFIREQDLYALKSVKIPV
						RNHGILMETHKELKPLLSPSSET
						TVTVELPEADRAGAGTGAQAG
						QLMGFFKGIDQDIERAVQSEIFL
						HESYCMDTSHQPLLPAPPKTPM
						DGADCGIQWWNAVFIMLLIGIV
						LPVFYLVYFKIQASGETPNSLNT
						TVIPNGSMAMGTVPGQAPRLA
						VAVPAVTSADSQFSQTTQAGS
20104	50472	Α	20220	1	626	RYSSTLTRPWTQTPRTRASPHS\
						PVAGALTARPSPPRPTKPSRAG
						VHASRRPPQDLGPIGKGSAKTK
						GPRHQRKHHESITPPSRSPTKEA
						PPGLNDDAQIPASQESVAPLWIP
						VTAP*AHRVLSSCEFGAAFESA
						GEEEGEGEVSASQAAVQA\GTQ
		İ				TRR*GATSAPCGPVPEGASVPA
						L*AAGAAGRAEAERPPCVPAQ
						AVGLSGQPLS

SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	1	Amino acid sequence ( X=Unknown,
of peptide	hod		location of first	•	*=Stop codon, /=possible nucleotide
sequence		09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
			sequence		
50473	Α	20221	2492	4619	WHSWRQQGWGAHLGSSATGP
					GC*QGRPTCVMRPAGT/TRPGE
					APIGSGTGAVEMPAREPGDEKP
	İ				PAPFP*QRHFVPRAHLLFGTAL
	ŀ				RSLPELMRRAVRNPRHHLSAEE
					SLLFSLPGSPWKGWSSWNGQA
					PPLAPPAQGPLLPVPLSAGPGV
					AGERTVPSSPRWWRPHLFLAPT
					APSPPPCPTVSAPGHLLGVAQK
			! !		PDCHPPASQRPRRHPLPFLPGEA
:					FAE/SAANS*ISSISASFTFS*SSS
					WSLD\STKSSLPPLHLPTLTSRV
					PGKFSALLSQFSASFSSHGLPSE
					SVAIGTCST/PGERAHSRQTCGP
					GSSAQS*GQEEGQCPQCRDSAG
					TCGRPRCSRHPAGHAGSEPRPL
			•		QRRSLHCSHTG*GAGIGHKART
					CGRKAPPRCPSETGHHSGVGY*
					PWRQSQQPAARGRARCGRPCP
					GACPAMAPQTQSEEGRRAPR/T
					NPAVPSEPARP/STAGPRAGASV
					PPSPGPRGPAGPGAPS\AGRRRR
					APGATACSGGAVESG*PSL/GSA
					LSAQVAPEAARAPA*GPGQAA
					GAREEGPAWPCRSPP\GPGPSQC
					AALPQK/GSALRSGGRRTLGPL
					LSHPSFKILLAGSGGVLDNSTSS
					ERSRATRSSICRPVVSESWTNN
	l				NGGDTGG\SSCDFYSPGPADGL
					SQGHSSDHTNLLLVLSSQKRHF
		}			PA*ALVQPHPAPSPTGTEPWPP
		į			AGPRTVPRPPSASSVAPSCWWV
50474	A	20222	103	538	PQLSLPLLSIRDRRRPSKCANLT
					LTLTVAARHGLQSLGKVGISRR
		ļ			MPPPANLPSLKAENKGNDPNV
	1				NIVPKDGTGWASKQEQHEEEK
					TPEVPPAQPKPGVAAPPEVAPA
					PKSWASNKQGGQGDGIQVK*Q
		}	1	}	FQQEFPSLQAAGDQK
50475	С	20223	1	569	
50476	A	20224	1	1095	-
	50473 50474	50473 A  50474 A	of peptide sequence hod in USSN 09/540,217  50473 A 20221  50474 A 20222  50475 C 20223	of peptide sequence         hod 109/540,217         location of first codon for peptide sequence           50473         A         20221         2492           50474         A         20222         103           50475         C         20223         1	of peptide sequence         hod in USSN 09/540,217         location of first codon for peptide sequence         codon for last amino acid of peptide sequence           50473         A         20221         2492         4619           50474         A         20222         103         538           50475         C         20223         1         569

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20109	50477	A	20225	1	1970	MWKGFTILDSFKNISDSWEDVR
						ISTLTGVWKKLIVTLMDDLEEF
ŀ						KTSVEEITLTQRKLFFVMESPV
						GPISPPQPPSVSAWNKPLTSFGS
ļ						APSSEGAKNGQESGLEIGTDTIQ
						FGAPASNGNENEVVPVLSEKSA
						DKIPEPKEQRQKQPRAGPIKAQ
						KLPDLSPVDNK\EHKPGPIGKER
						SLKNRKVKDAQQVEPEGQEKP
						SPATVRSTDPVTTKETKAVSEM
						STEIGTMISVSSAEYGTNAKESV
						TDYTTPSSSLPNTVATNNTKME
						DTLVNNMESARKAWENSPNVR
]						EKGSPVTSTAPPIATGVSSSASG
						PSTANYNSFSSASMPQIPVASVT
						PTASLSGAGTYTTSSLSTKSTTT
						SDPPNICKVKPQQLQTSSLPSAS
						HFSQLSCMPSLIAQQQQNPQVY
						VSQSAA/ATLTTACLS*WSTAQI
						PAFYMDTSHLFNTQHARLAPPS
		1				LAQQQGFQPGLS\RSQPAFMQS
		1				SLSQPSVVLSGTAIHNFPTVQH
						QELAKAQSGLAFQQTSNTQPIPI
						LYEHQLGQASGLGGSQLIDTHL
						LQVTVPLPASQLSLPNFGSTGQ
						PLIALPQTLQPPLQHTTPQAQA
		:				QSLSRPAQVSQPFRGLIPAGTQ
		ŀ				
						HSMIATTGKMSEMELKAFGSGI
		ŀ				DIKPGTPPIAGRSTTPTSSP\SGL LLQVRTASPAK
20110	50478	<u> </u>	20226	260	8485	VSTEMSEKSGQSTKAKDGKKY
20110	304/8	A	20220	200	0403	ATLS\LFNTSKGKSLETQKTTVA
						ARHGLQSLGKVGISR\RMPPPC*
						PPKFLKPE*QRANDPNVNIVPK
		1				,
						DGTGWASKQEQHEEEK\TPEVP
				1		PA\QPKPGVAAPPEVAPAPKSW
				1		ASNKQGGQGDGIQVNSQFQQE FPSLQAAVDQEKK\EKETNDDN
						1
						YGP*PSLRPPNVACWRDGGKA
				1		AGSPSSSDQDEKLPGQDESTAG
						TSEQNDILKVVEKRIACGPPQA
<u></u>		1_				KLNGQQAALASQYR
20111	50479	Α	20227	330	485	RQGDRVSLCCLGRS*TPGLK*F
		1				SHLDLLNCWDYRCETVHLAEI
		<u> </u>				AQVSEKQI
20112	50480	C	20228	1189	1413	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	1	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
20112	50491	A	20229	<u>L</u>	407	MPFSLADCDSQHCLLFAVVDL
20113	50481	A	20229	1	407	GTAKATSCMSDTFDFYSSSGNS
						ANLKEGGKSPPAGPGKGPRGA
		1	ļ			QKRKYSGPWNRKESCKKHEGV
						KGAEGRGGTVASVRES/REPRF
						KMAAFSMHPC*RLSRHRGHLA
						QRCFCFCR
20114	50482	A	20230	1	408	HMDLAQPSQPVDELELSVLERQ
20114	30462	^	20230		400	PEENTPLNGADEVFPSLDEEVPP
						AEANKESPWSSCNKNVVGRCK
						LWMIITSIFLGVITVIIIGLCLAA
						VTYVDEDENEILELSSNKTFFIM
		1				LKIPEECVAEEELPHLLTERL**S
		1				L**HLGKWR**SST/GLHLPTTF
		1	1	<b>.</b>		LLQELQGLSLLASAGGTSSSKE
			į			GKTSSAPLRGVFSSGCRSSTESS
						SSSTGCEGWARSM
20115	50483	С	20231	144	383	ossi debe wines.
20116	50484	A	20232	11	530	AKDSESRRMLRGYNPQQMSEG
20110	30404	1	20232	1		HROKSEFYNRTASESNVYLNSF
					,	HYPDHSYKDQAFDTLSLDSSDS
						METSISACSPDNIS\SASTSNIARI
						EEMERLLKQAHAEKTRLLESRE
						REMEAKKRALEEEKRRREILEK
						RLQEETSQRQKLIEKEVKIRERQ
						RAQARPLTRYLPVRKEDFDL
20117	50485	В	20233	10	543	
20118	50486	A	20234	370	556	
20119	50487	Α	20235	3	3160	TELQQLGQKTADGSPGLKRALI
1						FALNFHSSPPGKDMAAVQRTL
						MALGSVAVTKDDGCYRGEPS
						WFHRKAQQNRRGFSEEQLRQG
				1		QNVIGLQMGSNKGASQAGMTG
						LSDPTPWTAHLGMYVQGNAKF
Ì						VSPVRRIVVHEYYNSQTFDYDI
				İ		ALLQLSIAWPETLKQLIQPICIPP
				ŀ		TGQRVRSGEKCWVTGWGRRH
						EADNKGSLVLQQAEVELIDQTL
				İ		CVSTYGIITSRMLCAGIMSGKR
	<u> </u>					DACKKRKLGTERPGTQAAP
20120	50488	Α	20236	231	873	KKFRALGSVGPGFRASGVQVA
						DEV\CRIF\YDMKVR\KCS\TPEE
						DQDKERSAVIF\CLSADKKCIIV
						E\EG\KEILVGDVGWNP*TDPF\R
						HFVG\MLP*KKIVRYAFVMDGK
						LLETKGFQKRKSLMFFFV/WAP
						EL\APLKK*K*SYCKLQGMAFK
						KNFQGIKHECQA\NGPEDLNRA\
						CIAEKL\GGSLIVAFEGCPCVDH
				<u> </u>		HSVPQIEKLPMFNCYPLAI

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20121	50489	A	20237	915	1255	GHGQKPAGKKKKILAVRSGSEP AGGFQGY/RGGHGTAS/GSPGD DDSPGALWEQSPGPL*AWPGA VSEEPMVCAEPVQEQSGPQLPP PRHSRSPSGDSGQPGVPAERRII MGMADA
20122	50490	A	20238	3	227	LLMDTF/FQIVIYL/GETIAQWR KAGYQDMPEYENFKHLLQAPL DDAQEILQARFPMPVKLRIQPG NATVSDCVQNV
20123	50491	A	20239		642	RRDTIARNWANVQSQVRHIEQ AFDQEAAA\VLMARLG\VFEPK SKEGPDVLRWLDRQLIRLCQKF GQYTKEDPTSFRLSDSFSLYPQS MFHLRRSPFLQVFNNSPDESSY YRHHFARQDLTQSLIMIQPILYS YSFHGPPEPVLLDSS\RFPMPRYI NTEHGGSQARFLLSKVNPSQTH NNLYAWGQETGAPILTDDVSL QVFMDHLKKLAVSSAC
20124	50492	В	20240	20	277	
20125	50493	Α	20241	1	3030	

2565

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20126	50494	Α	20242	1	2677	MAEVKVKVQPPDADPVEIENRI
						IELCHQFPHGITDQVIQNEMPHI
						EAQQRAVAINRLLSMGQLDLL
						RSNTGLLYRIKDSQNAGKMKG
						SDNQEKLVYQIIEDAGNKGIWS
						RDIRYKSNLPLTEINKILKNLES
						KKLIKAVKSVALGIPAGCSVCP
				:		KCGASNHLSARFCGSCGICVKS
1						LVKLSLDRSLALAAEEPRPFSES
						LNIPLPRSDVGTKRDIGTQTVGL
						FYPSGKLLAKKEQELASQKQR
						QEKMSDHKPLLTAISPGRENIG
						ENLQVIGLVKNFLSNIPQAPAT
1						QAKMGKWDPRQRGAQSPLIFL
						YVVDTCLEEDDLQALKESLQM
						SLSLLPP\DALVGLITFGRMVQV
				i		HELSCEGISKSYVFRGTKDLTA
						KQIQDMLGLTKPAMPMQQARP
						\AQPQEHPFASSRFLQP\VHKID
						MNLTDLLGELQRDPWPVTQGK
			•			RPLRSTGVALSI\AVGLLEGTFP
						NTG\ARIMLFTGGPPTQGPGMV
						VGDELKIPI\RSWHDI\EKDNARF
						\MKKATKHYEMLANRTAANGH
						CIDIYACALDQTGLLEMKCCAN
						LTGGYMVMGDSFNTSLFKQTF
						QRIFTKDFNGDFRMAFGATLDV
			:			KRVQVGERGREAKNKLWTVH
						AHSSWRALNTGLRSFDFIPGTV
						REPVKILGRRLTSDLQIRNSPLE
			i			AAVQGGVSKTVSTVSLAHAMR
		L				AHPHLLMEDLRILVIDLDPQSS
20127	50495	В	20243	453	3318	
20128	50496	Α	20244	2	1367	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
20129	50497	Α	20245	1	1459	MLEGKYGPPPGSFQTRRTALAR
		1				SHTNTSPSTSPAANRLNALGIGE
						RDASPSWLGETQDQVKRQPGL
		l				SGSRKIPRCAMTYGSLGQQAA
		1				QAAGVHAPNVFVASLLCQFLQ
		1		ļ		FVYRNSMGWTGGADFSYSCQE
		1				GETKELWSSGHHLKACRAVAF
		1				SEDGQSEYWGRQPAMWWKGA
		ļ				VSSTMTLGTFSPELITVSKDKAI
						HVLDVEQGQLERRVSKAHGAP
		ļ				INSLLLVDENVLATGDDTGGIC
		ļ				LWDQRKEGPLMDMRQHEEYIA
		ŀ				DMALDPAKKLLLTA\TYCSTLY
						SGDGCLGIFNIKRRRFELLSEPQ
						SGDLTSVTLMKVQLVMWGKK
						VACGSSEGTIYLFNWNGFGATS
		İ				DRFALRAESIDCMVPVTESLLC
					ļ	TGSTDGVIRAVNILPNRVVGSV
				,		GQHIGEPVEELALSHCGRFLAS
						SGHDQRLKFWDMAQLASCGG
						G*LPSAQKKGRTTAGSEQQDLE
						HR*LLRRTEARGRSLHGSQKKR
	İ					RRLGMTVTEGMN
20130	50498	Α	20246	221	847	RHIFSFFSGHVLAVSLGKFIFEST
						LFPSERAARRRRRPVQQWPRFK
						ARDLRGKKKEELLKQLD\DLKV
1		İ				EAV/SQLRVAKVTGR\AASKLS
		İ				KIR\VVRKSIARVLTVI\NQTQKE
						NLRKFYKGKKYKPL\DLR\P*GR
		İ				HRAMRPPASNQARRENLEGPK
						EAGSGKGAAFTRCGK*RGSRA
						LRGRLCQ*KHKENLKTKKQQR
		<u> </u>				KERLYPLRKYAVKA
20131	50499	Α	20247	1	415	FFFFFECG*SFTNSFSVTIHRRIH
						NGEKSYECSDCGKSFNVLSSVK
						KHMRTHTGKKPYECNYCGKSF
						TSNSYLSVHTRMHN/TANVNSI
						TVGKAFIDLSCLR*HEQTLTRC
						MNHLLLCNKLPPKILWLQTKTL
						VISKFL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20132	50500	Α	20248	196	1768	RRSVLLPGCPVAVGVTGWRLSS
				1		CPQLPLCLPCSQPLSEKDTQRA
						ESWLMSS*KAG/YKGLVTFEDV
						AVEFTQEEWALLDPAQRTLYR
				1		DVMLENCRNLASLGNQVDKPR
		1		ļ.		LISQLEQEDKVMTEERGILSGTC
		İ				PDVENPFKAKGLTPKLHVFRKE
						QSRNMKMERNHLGATLNECN
		1				QCFKVFSTKSSLTRHRKYSILGE
		1				RPYGCSECGKSYSSRSYLAVHK
		1				RIHNGVKPYECNNCGKTFSSRS
ļ		1				YLTVHKRIHNGEKPYDCSYCG
			1			KTFSNS*YLRPHLRIHTGEKPYK
		1				CNQCFREFRNQSIFTRHKRVHT
						GEGHYVCNQCGKAFGTRSSLSS
}						HYSIHTGEYPYECHDCGRTFRR
				ŀ		RSNLTQHIRTHTGEKPYTCNEC
						GKSFTNSFCLTIHRRIHNGEKSY
				<u> </u>		ECSDCGKSFNVLSSVKKHMRT
		İ				HTGKKPYECNYCGKSFTIGQGD
						GFEIVLPPPQLQQLIKAFFLGNT
						HCLSDRLSVQCIHYSIRYLHLNI
						HYLHYSIHYLHYSICYLHYSIM
						DPPQQSPHPLQHPLQNPLQ
20133	50501	Α	20249	2	408	
20134	50502	Α	20250	1	280	
20135	50503	Α	20251	2	469	RGIREGEEFDVDLAGMEVAVG
						GWGRGLWLHRHQAGIRGQSLG
		1				WRCRPWVGAGGVQAASERPD
						LAGSIMGPEVGALRRASPVIVIQ
						SQAGAFLSSNTTCQQFRDPGFR
						GRYPASPHTLSLPPETAHL*LLL
						CLPAAGVLPHTWPLLDRQGWH
	1		1	ļ		RGHAR

SEQ ID		1	SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20136	50504	Α	20252	1	1762	FRMAAPLVLVLVVAVTVRAAL
						FRSSLAEFISERVEVVSPLSSWK
						RVVEGLSLLDLGVSPYSGAVFH
	1					ETPLIIYLFHFLIDYAELVFMITD
						ALTAIALYFAIQDFNKVVFKKQ
						KLLLELDQYAPDVAELIRTPME
						MRYIPLKAALFYLLNPYTILSCV
						AKSTCAINNTLIAFFILTTIKGKA
						LKGEYKGSAFLSAIFLALATYQ
						SLYPLTLFVPGLLYLLQRQYIPV
	1					KMKSKAFWIFSWEYAMMYVG
Ī						SLVVIICLSFFLLSSWDFIPAVYG
İ						FILSVPDLTPNIG\LFWYFFAEM
						FEHFSLFFVCVFQINVFFYTIPLA
						4
						IKLKEHPIFFMFIQIAVIAIFKSYP
						TVGDVALYMAFFPVWNHLYRF
		1				LRNIFVLTCIIIVCSLLFPVLWHL
1						WIYAGSANSNFF*GITLTFNVG
i		1				QLPHELFIMVPEGGGPGALMA
						AIERNVDRAVPGGSAERDPIER
	<u> </u>					STLHILLISDYFYAFLRREYYLT
						HGLYLTPQGIAPEAMLVLQSRP
						GWHRASWTLRGLVGPEAGPKL
						SSPELPAGECLGRRGSSPRVTSS
						LVPKRDPWLTDSKAYGEELGA
	<u> </u>					PQLGPPPCGSAHQGAPSQTGRR
20137	50505	В	20253	102	406	
20138	50506	В	20254	200	408	
20139	50507	В	20255	347	458	
20140	50508	A	20256	2	430	
20141	50509	A	20257	3	567	
20142	50510	В	20258	1	1317	
20143	50511	A	20259	928	2067	
20144	50512	A	20260	2	4498	VHELQSLSEMLKPEVLSPTVKG
ł	[					KLLTGIKSPRSYTPKPNPVTHVI
						STGIQKIWIGATVGSTEVSVAV
	1					TPDGYADAVRGDRFMMPAERR
						LPLSFVLDVLEGRAQHPGVLYV
						QKQCSNLPSELPQLLPDLESHV
						PWASEALVHKHHYENLYCVVS
						GEKHFLFHPPSDRPFIPYELYTP
						ATYQLTEEGTFKVVDEEAMEK
						VSVLFLGSREGEGQKPGSEEQA
						QKIWGVEAAVQRLGMLEAGIL
						PQEGRFQALAEIADI
L	I	<u> </u>	l			- x x x.

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20145	50513	Α	20261	3	3137	DTPAWHGDANRTKAALEAARS
						DYLEFPPAARELCVPLAVPYLD
						KPPTPLHFYRDWVCPNRPCIIRN
						ALQHWPALQKWSLPYFRATVG
						STKVSV\AVTPDGYADAVERGI
						RF/M*LPVERRLPLSFVLDVLEG
						RAQHPGVLYVQKQCSNLPSELP
						QLLPDLESHVPWASEALGKMP
			i			DAVNFWLGEAV*VTFLHKDHY
						ENLYCVGSGEKHFLFHPPSDRP
						FIPYELYTPATYQLTEEGTFKVV
						DEEAMEKAEVSRTC
20146	50514	Α	20262	164	1236	AQSRRSRDRGQAFSNHHAAEHI
						TCTPGSGSGSRVSSGQAARRRL
						SGRAGLWAAVTNPPPHPTWLE
			:			VSRSAAATARAALCRRSACLLP
	}					RTPHLSAMISASRAAAARL/VG
						AAASRGLRAARHQIRSNQGTV
						VGIDLGTTNSCVAVMKKNVPF
						KIVRASNGDAW/VEAHGKLYSP
1						SQIGAFV/LMKMKETAD**LFLS
						SEPPQPGQRAGFLQPSCCGAYY
1						LYPWLRERQSSILWSGGAAAPQ
!						RKSGPLGRSDQPPPSPHVVGGF
						QKRCRHRSRSSLPSERLFAASYS
1						SFIRHDKCQPSCSSPSRGAAASR
						GLRAARHQIRSNQGTVVGIDLG
}						TTNSCVAVMKKNVPFKIVRAS
						NGDAWVRLMGNCILRVRLEHL
						CDEDERDCRLIVALPEGMEEPE
ŀ						FDYQNDMGNSQDLERRRGCAA
						EPWEEGVGQSLALPLVSLCLKC
						RWYRSPCTQGQHPATFSGPAEG
						VISERSMLEHWHEEINQCKAYN
						EDVFSTMRTNPARISSEAMAPA
						DIIDHSSQRPDFTSGKMLADEF
						GTASNIKSRVNAFQSWEPLHLY
						NKDSNL
20147	50515	Α	20263	216	728	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	1 ' ' '
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20148	50516	Α	20264	<u> </u>	2519	MAPSSRSGAGNTSTPGGCARAR
20146	130310	Α.	20204	. 1	2317	APPGNAGVHSGTQTGAVRDTA
						FISPSSPGLFLSSEPPQPGQRAGF
						LQPSCCGAYYLYPWLRERQSSI
İ						LWSGGAAAPQRKSGPLGRSDQ
						PPPSPHVVGGFQKRCRHRIAQL
İ		İ				FAVGALVCCLVLLHLSAMISAS
1						
l						RAAAARLVGAAASRGPT\AAR
						HQDSW\NGLSHEAFRLVSRRDY
			İ			\AS\EAIKGAVVGI\DLGTTNSCV
			İ			AVMEGNQAK\VLENAEGAKTT
						PSVVAFTADGERLGGMPAKRQ
ŀ						AVTNPNNTFYATKRLIGRRYDD
	1					PEVQKDIKNVPFKIVRASNGDA
				1		WVEAHGKLYSPSQIGAFVLMK
1						MKETAENYLGHTAKNAVITVP
						AYFNDSQRQATKDAGQISGLN
						VLRVINEPTAAALAYGLDKSED
1	1					KVIAVYDLGGGTFDISILEIQKG
		1				VFEVKSTNGDTFLGGEDFDQAL
ľ		1				LRHIVKEFKRETGVDLTKDNM
						ALQRVREAAEKAKCELSSSVQT
	ŀ	]				DINLPYLTMDSSGPKHLNMKLT
		1				RAQFEGIVTDLIRRTIAPCQKA
						MQDAEVSKSDIGEVILVGGMT
1						RMPKVQQTVQDLFGRAPSKAV
1						NPDEAVAIGAAIQGGVLAGDV
						TDVLLLDVTPLSLGIET\HGAVF
						SKLINRKTNIPTKKSQVFSTAAD
	1	1				GQTQVEIKVCQGEREMAGDNK
						LLGQFTLIGIPPAPRGVPQIEVTF
			<u> </u>			DIDANGIVHVSAKDKGTGREQ
20149	50517	A	20265	1	168	LERERKKSRKELLNFYAWQH
20179	150517	~	20203	[ '	1.00	RESKMEHLR/QLRKKFEEDKQR
						IELLRAQRKFRPY
I .	I	L				IELLKAYKKEKEI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20150	50518	A	20266	3	1085	ARAGAEPQLPATPALPGGKMV
						ARSLAGLSSG*PGHPSPPISFPAI
						PIKFSEKQQASHYLYVRAHGIQ
						QGTKWREALGWP/SRMLTLGF
						P/PPCS/EESLSCLQSTCGLVQSV
	j					ELKFPDLPHPCTSCLAFSVPC*S
						SRTVSAGHLQGLPRLPSCFPTE
						GAPSPGLGAGSPGALLFRDLVP
						DVGRSPQPLTSLSLQGFQVAYV
						VFQKPSGVSAALAVGPKCQA*
		İ				GSPVSLAPFSSEWISDYADSVPD
				1		PGWRLREVPGAHRTMGSPSLQ
						EEAKAKEEEGVPDEEGWVKVT
				1		RRGRRPVLPRTEAASLRVLERE
						RRKRSRK\ELL\NFYAWQHSRE/
						SKMEHS*GKCARSFEEDKQRIR
		1				ACCGPKRKIPDRN
20151	50519	Α	20267	2	334	SLQVQPFASLISKAPLSTPRLLIN
1						KEKAGQSDPFLGMIMGLGGGM
						DFDSKKA/YAELLGWKKELEDL
ŀ						VRREHASIDAQSGAGVPNPSTS
	}					ASPKKSPPPAKDEARTTEREKP
						Q
20152	50520	Α	20268	124	407	QSDPFLGMIMGLGGGMDFDSK
1		1				KAY/RVRPRRGLGQPGMDRVR
			ļ			WERGHEGR*EEQAGASPPGNS
ļ		1				DLLLHRRDVAWLGECDQGCLA
		<u> </u>	ļ			LAELLGWKVRS
20153	50521	A	20269	193	458	DDD AVCDCL V CECTA DIA A CDD
20154	50522	Α	20270	55	1165	RDRAVGDSLKGFSTAPMAEPD
		1				PSHPLETQAGKVQEAQDSDSDS EGGAAGGEADMDFLRNLFSQT
						-
				1		LSLG\SQKERLLDELTLEGVA\R
1						YMQSERCSQSSSCLVGAGISTS AGIPRLFASPSTGLLLNNLEK*P
						SFPNPEAIF*DSAIFKKH\PEPFFA
						LAK\EL\YPGAVSRPTICSIISCGL
ļ.						
						LERDKGGLLLR/CVKRREH*VT
ľ						WERIAG\LEQEDLVEAHGTFYT
						SHCVSASCRHEYPLSWMKEKIF
						SEVTPKCEDC\QSLVKPD\IVFFG
						ESLGARFFSCYCSSDFLKV\DLL
						L\VMGTSL\RDVAWLGECDQGC
				1		LALAELLGWKKELEDLVRREH
						ASIDAQSGAGVPNPSTSASPKK
L			1	<u> </u>	<u> </u>	SPPPAKDEARTTEREKPQ

	of peptide sequence	hod	in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20155	50523	A	20271	3	357	FGFNGCSKRIIKLQELSDLEERE NEDSMVPLPKQSLKFFCALEVV LPSCDCRSPGIGLVEEPMDKVE EGPLSFLMKRKTAQKLAIQKAL SDAFQKLLIVVLG/QDCLDHP*S TSVSVSK
20156	50524	Α	20272	696	1123	IPLKMMNFLQSLKSPQLSHQLR L*LPQEVEPSLTQQNFPAQTP/V PPMEAEPSPNQQEATVQASEPP KNIELSSQQMVPENIFPPTMENS NQLPEPPTEVVAQLPPRYEVTIP TQGQDQAQLSTLASVTLQPLDL GFIITPESTTEIELSPTMQETPTQ PPKEFVPQPPV/LSRESSEETGPL PVQQETSAESPEPTKDENPSPIQ *EAAFSEISSQGSSWHQT
20157	50525	A	20273	1	727	GIPAADR\EASLELIKLDISRTFP NLCIFQQGGPYHDMLHSILGAY TCYRPDVGYVQGMSFIAAVLIL NLDTADAFIAFSNLLNKPCQMA FFRVDHGLMLTYFAAFEVFFEE NLPKLFAHFKKNNLTPDIYLID WIFTLYSKSLPLDLACRIWDVF CRDGEEFLFRTALGILKLFEDIL TKMDFIHMAQFLTRLPEDLPAE ELFASIATIQMQSRNKKWAQVL TALQKDSREMEKGSPSLRH
20158	50526	A	20274	3	435	CGGIPKMEESAEYEAARHKRE KRMENINLAGSKRGKEDGKDS EFAAIENLPGFELLSDREKVLCS SLNLSPARYVTVKTIIIKDHLQK RQGIP/CQKPPS*LPGQSPKEKD FEFPHRKRLDLQGRVLKLRRFE SQGDAQTVCQPK

SEQ ID NO:	SEQ ID NO: of peptide	Met	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last	Amino acid sequence ( X=Unknown,
	sequence		09/540,217	codon for peptide sequence	I .	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20160	50528	A	20276	3	1084	NQEWKRSPPITVAKATQGAYL
		İ				NILKDTLRAQKEQKEGKEEVIH
ĺ		1				KGAKLHPNGYRMAQGSETLVA
						RGGPCRSVEPSAASPQELRGW
ľ						WEAQALKRWGLMGGVWVME
		1				VDPSWLGAISAIVSSSASRLLKS
	]					VWHIPCPHFLLLRPQLKEAQRR
						KKQLEERCRVEESIGNAVLTW
				1		NNEILPNWETMWCSRKVRDLW
						WQGIPPSVRGKVWSLAIGNELN
		ŀ				ITHELFDICLARAKERWRSLSTO
		1				GSEVENEGLCVAVCAQQGHVG
						VMGFGSDEPSAVSPCEKGKSLA
						AWVLIFVDFRVGLQKSFQKRK
						ERESTKLQQLWSWCLMLTYFA
	İ					AFEVFFEENLPKLFAHFKKNNL
						TPDIYLID/W*FRLLVGC
20161	50529	Α	20277	60	556	EDSEMAPPAPGPASGGFGEVYE
						LFDVKNAFYIGSYQQCINEAQR
						VKLSSPERDVERDVFLYRAYL\
					RRDSIVAELDREMSRSVDVTNT	
						TFLLMAASIYLHDQNPDAALRA
						LHQGDSLECTAMTVQILLKLDR
						LDLARKELKRMQDLDEDATLT
20162	50500	[				QLATAWVSLATGG
20162	50530	A	20278	1		FFYKSVLAQKEVRCVLDE/ISPL
						GPPELKPVRMFADL/LPHESRR
					1	DSSVPELAEKMSRSVDVPTPPS
						LLMAASIYLHDQNPDAALRAL
						HQGDSLECTAMTMQDAYYIFQ
						EMADKCSPTLLLLNGQAACHM
						AQGRWEAAEGLLQEALDKDSG
						YPETLVNLIVLSQHLGKPPEVT
						NRYLSQLKDAHRSHPFIKEYQA
20163	50531	A :	20279	1		KENDFDRLVLQYAPSA
20103	50551	^	20219	1		RKFGVVLDEIKPSSAPELQAVR\
				ľ	į į	MFADYLAHESRRDSIVAELDRE
						MSRSVDVTNTTFLLMAASIYLH
						DQNPDAALRALHQGDSL\ECTA
			ł			MTVQILLK\LDRL\DLARKELKR
	ŀ	1				MQDL\DEDATLTQLATAW\VSL
1		ŀ				ATGGEKLQDAYYIFQEMADKC
			ľ			SPTLLLLNGQAACHMAQGRWE
						AAEGLLQEALDKPPHGCSCERK
						LVSPFMVETEAQKSPVCKKVPC
l					L.	HGPPWAGRALGASSLDSGYPE
j						LVNLIVLSQHLGKPPEVTNRY
l	ļ					LSQLKDAHRSHPFIKEYQAKEN
0164	50532 I	3 2	20280	512	903	DFDRLVLQYAPSA
					413	
3.00	1	• 12	.0201		†1J	

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
20166	50534	A	20282	11	1195	MKHPNINSALQAPHPFVPGRPH
						LVAGVAPPALSNHPQARATAA
			•			AGKPHPRCAEAAGIGLACTAPG
						SHWLKMGELPLDINIQEPRWD
						QSTFLGRAP\HFFTVTDPRNLLL
				ļ		SGAQLEASRNIVQNYRAGVVTP
				ļ		GITEDQLWRAKYVYDSAFHPD
				ļ		TGEKVVLIGRMSAQVPMNMTI
						TGCMLTFYRQGSKDEGHCRRG
						RSECLCSLRKTPTVVFWQWVN
				-		QSFNAIVNYSNRRW*HFPSLVR
Ì						QLGT\AYVSAT\TGAVATALGL
						KSLTKHLPPLVGRFVPFAAVAA
						ANCINIPLMRQRELQVGIPVAD
						EAGQRLGYSVTAAKQGIFQVVI
						SRICMAIPAMAIPPLIMDTLEKK
						DFLKRRPWLGAPLQVGLVGFC
		ŀ				SIHISNLEPELRAQIHEQNPSVE
2016	50505	_	20202		7.1	VVYYNKGL
20167	50535	В	20283	24	51	
20168	50536	C	20284	1	1182   372	
20169 20170	50537	B A	20285	1	4610	ELAQNHSRPAPASAPGGGGG
20170	20170   50538	I <sup>A</sup>	20200	1	4010	DPNMATARTFGPEREAEPAKE
	ļ					ARVVGSELVDTYTVYIIQVTDG
1						SHEWTVKHRYSDFHDLHEKLV
						AERKIDKNLLPPKKIIGKNSRSL
						VEKREKDLEVYLQKLLAAFPG
	İ					VTPRVLAHFLHFHF\YET*VIHA
ŀ						A\LAEELFEKGEQLLGAGEVFAI
						GPLQLYAVTEQLQQGKPTCAS
ŀ						GDAKTDLGHIL\DFTCRLKYLK
						VSGTEGPFGTSNIQEQLLPFDLS
						IFKSLHQVEISHCD
20171	50539	Α	20287	1	4670	MAAQRAAGATGGGGRGASAP
				ŀ		GGGGGGDPNMATARTFGPERE
				1		AEPAKEARVVGSELVDTYTVYI
						IQVTDGSHEWTVKHRYSDFHD
				1	;	LHEKLVAERKIDKNLLPPKKIIG
						KNSRSLVEKREKDLEVYLQKLL
į						AAFPGVTPRVLAHFLHFHFYEI
						NGITAALAEELFEKGEQLLGAG
l						EVFAIGPLQLYAVTEQLQQGKP
						TCASGDAKTDLGHILDFTCRLK
				1		YLKVSGTEGPFGTSNIQEQLLPF
		L				DLSIFKSLHQVEISH
20172	50540	Α	20288	2	227	ERFRPPIARVIDVSNGKVHVAE/
						SCLEETGGLGVDIVLDAG\LDPP
						DSHCLFLKGATLAFLNDEVWN
20:==	150517	<u> </u>	00000	ļ	107	LSNVQQGRYL
20173	50541	A	20289	3	487	<u></u>

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20174	50542	Α	20290	2	222	
20175	50543	A	20291	1	296	HLSPGKSV\LIMDGASA\F\GTIAI
						QLAHHRGAKVISTACSLEDKQC
						LERFRPPIARVIDVSNGKVHVA
						ESCLEETGGLGVDIVLDAGGYI
						FQHSSYIFT
20176	50544	Α	20292	353	512	
20177	50545	Α	20293	2	396	VRLYSKDDEPAVKLQLLPHKH
						DIITLLGVGGHWVTTEENLQLD
						PPDSHCLFLKGATLAFLNDEVW
						NLSNVQQGKYLS*QSFQLRILK
						DVMEKLSTGVFRPQSDEHIPVF
						EARI\SLEIVEKNQGR*NKLHF
20178	50546	В	20294	1	1182	
20179	50547	Α	20295	1 	1189	PTRPDQWEQGAGSLTKDWKV
						AVVKVQAVGAAQRQVTMKGL
						YFQQSSTDEEITFVFQEKEDLPV
						TEDNFVKLQVKACALSQINTKL
						LAEMKMKKDLFPVGREIA\GIV
						LDVGSKVSFFQP\DDEVVGILP\
						LDS*KTPGL\CESLLRVHEPLLG
						FINPEKVTW\TEAAGSISGMGV
						RAYYSFLHLS/YLHLLSWGKSV
l						LIMGGS/ASAFGTIAIQLAHHRG
						AKVISTACSLEDKQCLERFRPPI
]						ARVIDVSNGKVHVAESCLEETG
}						GLGVDIVLDAGVRLYSKDDEP
						AVKLQLLPHKHDIITLLG\VGGH
						WVTTEENLQLDPPDSHCLF\LK
						GATLAFLNDEVWNLSNVQQGK
						YLCILKDVMEKLSTGVFRPQLD
						EPIPLYEAKVSMEAV\QKNQGR
20180	50548	A	20296	3	115	FFKFSFFNGC*SIKRQSQGQALD
						RDIICFVDICTKDC

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20181	50549	A	20297		1401	MKEEELTRQIVLVQEAGKGVV RTKDGACAEVWQALEELEKGL CAGIVRVEGMAAAPPSYCFVAF PPRAKDGLVVFGKNSARPRDE VQEVVYFSAADHEPESKVECT YISIDQVPRTYAIMISRPAWLW GAEMGANEHG\VCIANEAINTR EPAAEIEALLGMDLVRL\GLER GETAKEALDVIVSLLEEHGQGG NYFEDANSCHSFQSAYLIVDRD EAWVLETIGKYWAAEKVTEGV RCICSQLSLTTKMDAEHPELRS YAQSQGWWTGEGEFNFSEVFS PVEDHLDCGAGKDSLEKQEESI TVQTMMNTLRDKASGVCIDSE FFLTTASGVSVLPQNRSSPCIHY FTGTPDPSRSIFKPFIFVDDVKL VPKTQSPCFGDDDPAKKEPRFQ EKPDRRHELYKAHEWARAIIES
						DQEQGRKLRSTMLELEKQGL* AMEEILTSSEPLDPAEVGDLFY
20182	50550	A	20298		356	QTPKIVADKDYSVTANS\KIVV VTAGVRQQEGESRLNLVQRNV NVFKFIIPQIVKYSPDCIIIVVSNP VDILTYVTWK/ARFRYLMAEKL GIHPSSCHGWILGAHGDSSVAV WSGVNLAG
20183	50551	Α	20299	1	322	
20184	50552	Α	20300	25	1086	
20185	50553	Α	20301	1	864	
20186	50554	Α	20302	13	190	

SEQ ID NO:	of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20187	50555	A	20303		1350	MEKPKRLRWLWGPAPSSLLAE PAPEETHAADFVFSARLLQRSP EPSLSWGGRALAEEQKAEGSSC GSAERLVAFAAKTRPYLSLCK MATLKEKLIAPVAEEEATVPNN KITVVGVGQVGMACAISILGKS RAMEHTTAGAPYTPRQQTCQC NMMSLADELALVDVLEDKLKG EMMDLQHGSLFLQTPKIVADK DYSVTANSKIVVVTAGVRQQE GESRLNLVQRNVNVFKFIIPQIV KYSPDCII\IVVSNPRDI\LTYGT WKLSGLPKHRVIGSGCNLDSAR FRYLMAEKLGIHPSSCHGWILG EHGDSSVAVWSGVNVAGVSLQ ELNPEMGTDNDSENWKEVHK MVVESA\YEV\IQA*KGYTH\WA IGLSVA/EILNESMLKNLSR\IHP VSTMVKGMYGIENEVFLSLP\CI \LNAGGLTSVINP\KVKDDEVAQ LKKSADTLWDI\QKDLKDL
20188	50556	В	20304	1337	1706	ERRONDTEWDIQROERDE
20189	50557	A	20305	178	344	
20190	50558	A	20306	2	297	TKMTVNNLHPRVTEEDIV/GQP MKCNLHMNGNVITSDQPILLRL SDSPSMKKESELPRRVNSASSS NPPAEVDPDTILKALFKSSGAS VTTQPTEFKIKL
20191	50559	В	20307	8	478	
20192	50560	С	20308	146	391	
20193	50561	Α	20309	3	735	HRQKDARFRIKGKVQDAREML NSRKQQTTVPQKPRQVADARE KISLKRSSPAAFINPPIGTVTPAL KLTKTIQVPQQKAMAPLHPHPA GMRINVVNNHQAKQNLY\DLD EDDDGIASV\PTKQMKFAASGG FL\HHM\AGLSSSKLSMSKALPL TKVVQNDAYTAPALPSSIRTKA LTNMSRTLVNKEEPPKELPAAE PVLSPLEGTKMTVNNLHPRVTE EDIVELFCVCGALKRARLVHPG VAEVG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
20194	50562	Α	20310	1	1763	MEIASFVGPNVNTVGNLRPAQS
						AAGDGDEVQSVHSQRQDCQAT
						GSLSNGENNEKNAFTTYQQET
]					:	KNTTTPQSVYFLGVNKMSSQH
						REGSRGTSPPLPVLACFELAVA
						RVGPRTPEVVRRSLAAEGAGSR
						GLDCFRLARRIKMADISLDELIR
						KRGA\AAKGRLNARPGVGGVR
						SRVGIQQGLLSQSTRTATFQQR
						FDARQKIGLSDARLKLGVKDA
						REKLLQKDARFRIKGKVQDAR
						RDV*TFRKQQTT\VPQKPRQV\A
	1					D\ARERIQLGRGRSPAAFLNPPI
	:					GTGTPALKLTTTIQRMHVDPVG
						WILTTECFQVPQQKAMAPLHP
						HPAGMRINVVNNHQAKQNLY
		ļ				DLDEDDDGIASVPTKQMKFAA
						SGGFLHHMAGLSSSKLSMS\KA
						LPLTKVVQNDAYTAPALPSSIR
						TKALTNMSRTLVNKEEPPKELP
	ļ					AAEPVLSPLEGNKMTV\NNLHP
						RVTEEDIV\EVFCVCGGL/RRRA
						RLVPSGG*PKVVFCEKRDDAIT
					ļ	AYKKYNNRCLDGQPMKCNL\H
						MNGNVITSDQPILLRLSDSP\SM
						KKESELPRRVNSASSSNPPAEV
						DPDTILKALFKSSGASVTTQPTE
20195	50563	Α	20311	1	915	
20196	50564	Α	20312	3	3238	LDTDDDLNSDDYEYEDEAKLVI
						FPDHYEIALPNIEELPALVSECA
						RLLARCAWLCVSLKLSTLLTDD
						TGLLTVRNKEEVLARSAKVTIA
						CDAVLSSKSPYRKQDPDTWEN
						ELPVSKYANNLTQLDNGVRIPP
						SGWKCARCDLRENLWLNLTDG
				Ī		SVLCGKWFFDSSGGNGHALEH
						YRDMG\YPLAVKLGTITPDGAD
						VYSFQEEEP\VLDPHLAKHLAH
						F\GID\MLHMHGTENGLQ\D\NDI
	1					KLRVSEWEVIQES

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20197	50565	A	20313	2	2716	AEFRGIPRRRRRQTPRSGSGSAR SAPVRAEAMQRRGALFGMPGG SGGRKMAAGDIGELLVPHMPTI RVPRSGDRVYKNECAFSYDSPN SEGGLYVCMNTFLAFGREHVE RHFRKTGQSVYMHLKRHAREK VRGASGGALPKRRNSKIFLDLD TDDDLNSDDYEYED\*AFLVIFP DHYEIALPNIEELPALVTIACDA VLSSKSPYRKQDPHTWENELPV SKYANNLTQLDNGVRIPPSGW KSARCDLRENLWLN
20198	50566	Α	20314	2	322	
20199	50567	A	20315	1285	3331	GAGAPRDWRLRRRGGAALCLK VLFDVRSFPREAAALTPHRLLQ FPRSSESPGDQMMAAARLLPVP AGPQPLSFQAKLTFEDVAVLLS QDEWDRLCPAQRGLYRNVMM ETYGNVVSLGLPGCKPDIIFHLE LRGISLGPVPEWG*EEPGPVE*L L/ENVETKGESQNTDLSPKPLIS EQTVILGKTPLGRIDQENNETK* SFCLSPNSVDHREVQVLSQSMP LTPHQAVPSGERPYMCVECGK CFGRSSHLLQHQRIHTGEKPYV CSVCGKAFSQSSVLSKHRRIHT GEKPYERNRACISLYSE\PHLAH LSLIHTGEKPH*CLECPKAFTQL SHLIQHQRIHTGERPYVCPLCG KAFNHSTVLRSHQRVHTGEKP HRCNECGKTFSVKRTLLQHQRI HTREKPYTCSECGKAFNDRSV LIQHHNVHTREKPYECSECGKT FSHRSTLMNHERIHTEEKPYAC YECGKAFVQHSHLIQHQRVHT GEKPYVCGECGHAFSARRSLIQ HERIHTGEKPFQCTECGKAFSL KATLIVHLRTHTGEKPYECNSC GKAFS\RTQLLIQHQRIHTGEKP YECGECGGAFNQHGHLIQHQK SAQKSCDPWLTQE/CPFSQKLH VEPTSSLQPKRSLC*LYRKLFFG DSVSQNNSRKKHLACCSCGKTS ETTCFIFLNHLEV/HVGE*SYNC
20200	50568	A	20316	1	319	
20201	50569	Α	20317	1	2433	
20202	50570	A	20318	2	293	GELAGTQCTRGACPHSP*PLHR Q/HTTVTHITAFDPDSTGQQVW QDLLQDGQLDSPTGDGGSDWE MVGREVVPVLGPGVGGLVISRP CFHVRDPYPWG

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20203	50571	A	20319		537	GGGTITRGWRGQFCRWQLNPG MYQHRTVIADQMRTLRSREV** HAQGHKFTVCLRREGQTVYQQ VLSLERPSVLRSWNWGLCGYF AFYHALYPRAWTVYQLPGQNV TLTCRQITPILPHDYQVRPPHLY LPLPSLPQDLPLLETAWPARRG SHACNPNTLGGKAGGSRDQEIE TILANT
20204	50572	С	20320	3	170	
20205	50573	Α	20321	278	1018	RVGVSRRPEVRNWATRDCRAQ ERMTHCLLHGMGRAGAASLTP KPMSLISAYCGGLWLAAVAVM VQMAALCEGTGQSRISFLLSSA GAKKPMRDCCGMVSSGSLSSL KAAERTPGALFFSSSTPGRYYN YDSSSRPQSRSVMSDQVCWTV GVPEGLWP*EKGDTEVREEEEQ PERKMGFPWRWEVFPTQHVVR ALQTIFELNVQAFAGGAMGAV NGMQPHGVPDKSSVQSDEVW VGVVYGLAATMIQE
20206	50574	Α	20322	3	627	

2581

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20207	50575	Α	20323	270	2945	NTSSFKLSPTTGPPGRYLTTTGA
						RIWNRLARYLVSEGSGLGAIEV
	1					MGTQDPGNMGTGVPASEQISC
	1					AKEDPQVYCPEETGGTKDVQV
	1					TDCKSPEDSRPPKETDCCNPED
						SGQLMVSYEGKAMGYQVPPFG
						WRICLAHEFTEKRKPFQANNVS
						LSYMIKHIGMGLRYLQWRDRK
				1		THVEKKTPFIDMNKSVPPIQSYF
				1		\VPLGGLRGGTSTRGWRGQFCR
						WQLNPGMYQHRTVIADQFTVC
		l				LR\REGQTVYQQVLSL\ERPSVL
						RSWNWGLCGYFAFYHALYPRA
						WTVYQLPGQNVTLTCRQITPIL
•		1				PHDYQDSSLPVGVFVWDVENE
						GDEALDVSIMFSMRN/GTGWW
		1				RRCPRGFVGMSPSVWSVGVQT
					,	VRGLLLHSSNPSKPLHDGLWLA
						RSSRAATTVNPH/CTAFDPG\RT
						GAA\VWQDLLQDGQLDSPTGQ
						RTPTQKGVGIAGAVCVSSKLRP
1						RGQCRLEFSLAWDMPRIMFGA
	1					KGQVHYRRYTRVFGAG\GDAP
						PALSHYALCRYAEWEERISAW
ŀ						QSP\VLDDRSLPAWYKSALFNE
						LYFLADGGTVWLEVLEDSLPEE
						\RAETSVTSAPTLRDYRRFG*LE
		ŀ				GQEY\RMYNTYDVHFYASFALI
						MLWPKLELSLQYDMALATLRE
						DLTRRRYLMSGVMGTCEKEGT
						FIPHDIGGPRWMEPWASAFNAY
						\LIHDTADW\KDLNLKFVACRFI
20208	50576	С	20324	123	359	
20209	50577	Α	20325	1	259	KPVGEKDTWSGMRTTGQLRPA
						HGVRLKANKDSLYKPILRQKK
						HFNSLHIPKALQKALPFKNKPK
	<u></u>					TQAKAGKVPKDRRR*TVSNITV
20210	50578	Α	20326	297	585	
20211	50579	Α	20327	182	374	PDFRIAAT*VVLDLDKSIKIVKK
						LKLTGFPYKIFKNTSFIKACISIY
						SHIYKSPYCLRKGMKYL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20212	50580	Α	20328	1	1267	MAANRKDFIHRRCGPELSQSEA
				•		VPQGLCRAERSSALAMLSGSVP
						RRTQLRPCKGTQRRCRRGGEA
						DSGGEAVGGGAAERRSAAARR
						RGGRRRGGGEAVGGGAAERRS
						AAARRRGGRRRRGGGEAGGG
						GAAEARRAAAARRA
						AAARRAAAARRAAAARRIAES
						GGEAHSRGEAHSGGA
						GAERRRPRLHSPALKTHPGQSS
				1		EGHVEYKLNHVEFEDQDDEAR
						VQYEGFRPGMYVRVEIENVPCE
						FVQNFDPLYPIILGGLGNSEGNV
						GHVQMRLKKHLWYKKILKSQ
						DPIIFSVGWRRFQTILLYYIEDH
						NGRQRLLKYTPQHIHCGAAFW
						DIVFMRT\WYLVSIPAFYNPVTS
						LLKPVGEKDTWSGMWTTGQL
						RLAHGVRLKTTKDSLYKNIHMI
						VETDTYSDHEEINTS
20213	50581	Α	20329	66	1414	KEELYGDFEDLETGDVHKGKS
						GPDTQNEDVEK\KEEIDPDEEES
						AKKKHLDKKRKLKEMFDVEY
					1	DEGESTYFDDLKGEMQKQAQV
						N/RHTRFPPRAPRHVQLNHAEF
						EDQDDEARVQYEGFRPGMYVC
						VEIENVPCEFV*NLDPRYPIILG
	:					GLGNSEGNVGYVQVGPFAAYL
ļ						VPEALWISPPSIILPSHPPQMRLK
						KHRWYKKILKSRDPIIFSVGWR
						RFQTIPLCYIEDHNGRQRLLKY
						TPQHVHRGAAFRSSSSSSSSSSS
						SSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSS
						IAATGVVLDLDKSIKVVKKLKL
						TGFPYKIFKNTSFIKGMFNSALE
						VAKFEDAVIRTVSGIRGQIKRA
	· ·					L*APEGAFQDSLEDKLRMSDIV
						FMRTWYPVSIPAFYNPVTSLLK
						PVGEKDTWSGMWTTGQLRLA
						HGVRLKTNKDSLYKVLVVCVL
						VEMKPVLYRQ

	0 5099	
20214   50582   A   20330   421		AKDQKKHRKKNSGPKAAKKK KRLLQDLQLGD\EEDAWKRNP KAF\AFQSAVWMARSFHRTQD LKTKKHHIPVVD\RTPLEPPPI\V VVVMGPPKVGKSTLI*CLI\RNF TRQKLTEI\RGPVTIVSGKKLRL\ TIIDCGCDINMMIDLG\EVADLV LMLIDASFGFEMETFEFLNICQV HGFPKIMGVLTHLDSFKHNKQL KKTKKRLKHRFWTEVYPGAKL FYLSGMVHGEYQNQEIHNLGH
20215 50583 A 20331 56	467	
20216 50584 A 20332 282	2 452	
20217 50585 A 20333 1	1356	
20218 50586 A 20334 80	1096	GLGVRVLRCPGVYRGPCTRAA YLGWLWTTLSPYCCLRSADMN PIVLVHGGGAGPISKDRKERVH HGMVRATTVGYGILREGGSAV DAVQGAVVALEDDPEFNAGCG SVLNTNGEVEMDASIMDGKDL SAGAVSAVQCIANPIKLARLVM ENFPHCFLTDQGAAHFAGAMG VPEIPGKKLVTERNKKRLEKEK HEKGAHKTDCQKNLGTVGAV ALDCKGNVAYATSTGGIVNKM VGRVGDSPSLGAGGYADHDF* AVSTTGHGE\SFLKVNLARLTLF HIEQGKTLEEAADPIG/CG*MKS RVKGLGGLIVVSKTGDWVAK WTSTSMPWAAAKDGKLPLRN
20219 50587 A 20335 1	1416	WISTOM WILLIAMS SIGN EXCIT
20220 50588 A 20336 3	277	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	delction, \=possible nucleotide insertion)
				sequence		
20221	50589	A	20337	235	2811	VKMEAPLRPAADILRRNPOOD
						YELVQRVGSGTYGDVYKARNV
						HTGELAAVKIIKLEPGDDFSLIQ
						QEIFMVKECKHCNIVAYFGSYL
						SREKLWICMEYCGGGSLQDIYH
						VTGPLSELQIAYVCRETLQGLA
		İ				YLHTKGKMHRDIKGANILLTD
						HGDVKLADFGVAAKITATIAKR
		ļ				KSFIGTPYWMAPEVAAVEKNG
						GYNQLCDIWAVGITAIELGELQ
						PPMFDLHPMRALFLMSKSNFQP
						PKLKDKTKWSSTFHNFVKIALT
						KNPKKRPTCLKDF*HHTF\VAQ
		İ				PGLSRALAVELLDKVNNPDNH
						AHYTEADDDDFEPHAIIRHTIRS
1						TNRNARAERTASEINFDKLQFE
-						PPLRKETEARDEMGLSSDPNFM
						LQWNPFV*WWQITGKSTSKR\A
						IPPPLPPKPRISSYPEDNFPDEEK
						ASTIKHCPDSESRAPQILRRQSS
						PSCGPVAETSSIGNGDGISKLMS
						ENTEGSAQAPQLPRKNDKRDFP
						KPAINGLPPTPKVLMGACFSKV
						FDGCPLKINCATSWIHPDTKDQ
						YIIFGTEDGIYTLNLNELHEATM
						EQLFPRKCTWLYVINNTLMSLS
						EGKTFQLYSHNLIALFEHAKKP
	1					GLGCPIFQTHRFPDRILPRKFAL
1						TTKIPDTKGCHK\CCIVRNPYTG
					}	HKYLCGALQSGIVLLQWYEPM
						QKFMLIKHFDFPLPSPLNVFEM
						LVIPEQEYPMVCVAISKGTESN
20222	50590	С	20338	365	389	
20223	50591	Ā	20339	1	350	FDVTELCGQWLSRGGEIEGFRL
						SAHCSCDSRDNTLQVDINGFTT
						GRRGDLATIHGMNRPFLLLMA
						TPLERAQHLQSS/RAPCCVPQAL
						EPLPIVYYVGRKPKVEQLSNMI
						VRSCKCS
20224	50592	A	20340	1	708	
20225	50593	В	20341	229	1249	
20226	50594	В	20342	49	375	
20227	50595	Α	20343	42	305	GARAHAPVLVQEEDLHGPHPL
						RPERL/PSVPSGRSSPPASR/PVG
1						PGG/DQLRPSPDRQQQVSGME
	1					WC/PWSGRSPGLSPTVGPRGGC
20000	50.50.5		2024		1000	HPTST
20228	50596	В	20344	8	1990 1512	
20229	50597	A	20345	1	1312	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleofide insertion)
20230	50598	Α	20346	1	1663	SGRDRRGRAWRKRNREPPTRG
				•		RSPDPQGPSARPLPSLGARISAA
						DRLSGLAARLIGNSRPVAGANA
						GVGPDLSHAGLVPTCARRLRR
İ						KNGLGRRRRRPPPSPPRPSLESS
	;					SRLTTLGTSQPLKFNFFPCDLRI
						FQTFNC\WGI\PVL*AKHR\ADR
						MRRFGRPF*TQESFDLALLEEV
Í						WSEHDFQYLRQKLSPTYPAAH
				8		HFRSGIIGSGLCVFSKHPIQELT
İ						QHIYTLNGYPYMIHHGDWFSG
						KAVGLLVLHLNGMELNAYVTH
1						LHAEYNRQKDIYLTHLVG\QA
1						WELAQFIHHTSKKADVVLLCG
İ						DLNMHPEDLGCCLLKEWTGLH
						DAYLETRDFKGSEEGNTMVPK
						NCYVSQQELKPFPFGVRIDYVL
						YKAVSGFYISCKSFETTTGFDPH
						SGTPLSDHEALMATLFVRHSPP
						QQNPSSTHGP\AERSPL/MCVCL
						KEALDGSLGLGMA\QARWWA\
						TFA\SYVIGLGL\LLLALLCVLA
ļ				:		AGGGAGEAAILLWTPSVGLVL
						WAGAFYLFHVQEVNGLYRAQ
1				ŀ		AELQHVLGRAREAQDLGPEPQ
	<u></u>					LYALL\LGQQEGDRTKEQ
20231	50599	Α	20347	175	544	RLRCQKGIPPSLRGRAWQYLSG
				İ		GKVKLQQNPGKFDQ\LDMSPG
	·					DPTWLDVIERDLHRQFP\FHEM
						FVS/RGGH/GQQDLFRVLDAYT
				-		LYRPEGSYCKTQAPIAAVLLMH
						MP\AEQAFWCLVQICEKYL
20232	50600	Α	20348	1	434	CAAGIRHELRRRARLGWRTAT
						ERMGRACPRPGKACREPGRA*
						ARALTPQPPTNSALSGLTRRPT
						ASPS\PHRQVRLDRGLAGRQGR
	[					AGGSTPGGAEAEGVQVAGHAQ
						AMVVAAARDACPPQHVPPKDS
L						APKDSAPQDLAPQVSAH
20233	50601	Α	20349	3	510	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20234	50602	Α	20350		1669	MAKSNGENGPRAPAAGESLSG TRESLAQGPDAATTDELSSLGS DSEANGFAERRIDKFGFIVGSQ GAEGASILGQTVPSPHGRVGEG PPIRSYTASSTGTGNRLEEVPLE VLRQRESKWLDMLNNWDKW MAKKHKKIRLRCQKGIPPSLRG RAWQYLSGGKVKLQQNPGKF DELDMSPGDPKWLDVIERDLH RQFPFHEMFVSRGGHGQQDLF RVLKAYTLYRPEEGYCQAQAPI AAVLLMHMPAEQAFWCLVQI
20235	50603	Α	20351	1	243	
20236	50604	Α	20352	70	382	
20237	50605	Α	20353	2	322	
20238	50606	A	20354	86	215	
20239	50607	В	20355	1	735	
20240	50608	A	20356	3	1104	MSSPQAPEDGQGCGDRGA\PPG DLRSVLVTTVLNLEPLDEDLFR GRHYWVPAKRLFGGQ\IVGQA M\VAAATSV\SEDVHV\HS\LHC YF/VFRAGDP/KSLPVLYQVERT RTGVEA\FRLRSVKGP*NMGKPI F\ICQ\ASFQQA\QPSP\MQHQFS MP/TLLPTTEELV*IVRTLFDQY LRDPNLQKRYPLALNRI\AAQE VPIEIKPVNPSPLRPAAEEWEPK QMFW\VRARGY\IGKSTPWKM\ HC\CV\AAYIS\DYAFLGTALAA LTQWPSTRCHFHGPPLD/HNPM W\FHAPFRAD\HWMLYECESPW AGG\SRGL\VHGRLWRQDGVLA VTCAQEGVIRVKPQVSESKL LRRQLQPLLPLPKSPPRSGAHL
						GSLFRSPQPSYPVRPPGSAGTPP PACLPPAQPCQGSPAMNLFRFL GDLSHLLAIILLLKIWKSRSCA GISGKSQVLFAVVFTARYL\DLF TNYISLYNTCMNV\VYIACSFTT VWLIY\TKFK\ATYDG\NH\DTFR SGSSRVV/VPQAILGVSWSNHD FTPL\EILW\TFSIYLGVSGPSLPQ LFHG*GKTGRGRETHSPATYLF CAKAFNPHALSLSNW\IWR\YHF \EGFFD\LIAIVAGPGPRQSSYCD FF\YLYYSPKFLKGERKFEFCRH RPRFLSISSPRAAAGRAGGKAG RKMKSFSHPRGDFFKNPTSLCS PIPPSCPGLKGNSARIPGLWGAQ RLGAVCSFLAF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20242	50610	Α	20358	356	1356	TPTTSGRTRKMWPRPGT*PP/A
1						NCSANINLTHQPWFQVLEPQFR
						QFLFYRHCRYFPMLLNHPEKCR
			:			GDVYLLVVVKSVITQHDRREAI
					:	RQTWARAAVRGWGPSAVRTLF
				}		LLGTASKQEERTHYQQLLAYE
			:	•		DALYGDILQWGFLDTFFNLTLK
					;	EIHFLKWLDIYCPHVPFIFKGDD
				}		DVFVNPTNLLEFLADRQPQENL
				ļ		FVGDVLQHARPIRRKDNKYYIP
						GALYGKASYPPYAGGGGFLMA
	i e					GSLARRLHHACDTLELYPIDDV
						FLGMCLEVLGVQPTAHEGFKTF
		ŀ				GISRNRNSRMNKEPCFFRAMLV
		l				VHKLLPPELLAMWGLVHSNLT
1						CSRKLQVL
20243	50611	Α	20359	221	1579	CCVDEGLEPTCFERTEDIGGVW
						RFKVSEIFLGLEQVVLLFQGESQ
						LMSCFSDFPMPEDFPNFLHNSK
		İ				LLEYFRIFAKKFDLLKYIQFQVL
						YFWGNGFLCISSAHI*IENTQSN
1						GKEQSAVFDAVMVCSGHHILP
İ						HIPLKSFPGETRWDSQLFGVGF
						QVLYM*FEGKRILVIGMGNSGS
1						DIAVELSKNAAQV*CSLLTMYL
						EGRKWGCHTGDWDSVFHTRFR
						SMLRNVLPRTAVKWMIEQQM
		l				NRWFNHENYGLEPQNK*SYFA
						FLMVYSLVSKVV*RCLKVKST
						VKELTETSAIFEDGTVEENIDVII
						FATGYSFSFPFLEDSLVKVENN
						MVSLYKYIFPAHLDKSTLACIG
						LIQPLGSIFPTAELQARWVTRVF
						KGLCSLPSERTMMMDIIKRNEK
						RIDLFGESQSQTLQTNYVDYLD
						ELALEIGAKPDFCSLLFKDPKLA
	10000	<u> </u>	-		000	VRLYFGP\CNSY
20244	50612	A	20360	3 292	570	
20243	50614	A	20362	2	511	RLRSGPLRLPGADSGSGPKAVC
20240	30014	\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\	20302	<b> </b>	J. 1	SPPFIVAPTGRGYCGDHESSFGA
						MEEPGVTPQPYLGLLLEELRRV
						SPGAMSVTWP/EGSREPPGEGSS
						RPALGSKPPWSEVPKPVLVCCP
						APAR\FEAVVRLVGRLSGFCVM
						EEDLGLWEGREKKLALMLSGLI
						EEKSKLLEKFSLVQKE
	1	1	L	<u>.                                    </u>	1	DEMORDED A CKE

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20247	50615	A	20363		3262	FLSDLEENFENNIFSFQVCDVVL NHAPDFRRVYLPYVTNQTYQE RTFQSLMNSNSNFREVLEKLES DPVCQRLSLKSFLILPFQRITRL KLLLQNILKRTQPGSSEEAEAT KAHHALEQLIRDCNNNVQSMR RTEELIYLSQKIEFECKIFPLISQS RWLVKSGELTALEFSASPGLRR KLNTRPVHLHLFNDCLLLSRPR ESVTGVAGQGTRGEGDEERGG SEREREGSEKLRWISALAMPRE ELDLLECYTAE
20248	50616	A	20364		2339	TGRGYCGDHESSFGAMEEPGA TPQPYLGLLLEELRRVVAALPE GMRPDSNLYGFPWELVICAAV VGFFAVLFFLWRSFRSVRSRLY VGREKKLALMLSGLIEEKSKLL EKFSLVQKEYEGYEVESSLKDA SFEKEATEAQSLEATCEKLNRS NSELEDEILCLEKELKEEKSKHS EQDELMADISKRIQSLEDESKSL KSQVAEAKMTFQIFQMNEERL KIAIKDALNENSQLQESQKQLL QEAEVWKEQVSELNKQKVTFE DSKVHAEQVLNDKESHIKTLTE RLLKMKDWAAMLGEDITDDD NLELEMNSESENGAYLDNPPK GALKKLIHAAKLNASLKTLEGE RNQIYIQLSEVDKTKEELTEHIK NLQTQQASLQSENTHFENENQ KLQQKLKVMTELYQENEMKL HRKLTVEENYRLEKEEKLSKV DEKISHATEELETYRKRAKDLE EELERTIHSYQGQIISHEKKAHD NWLAARNAERNLNDLRKENA HNRQKLTETELKFELLEKDPYA LDVPNTAFGRGSRGPGNPLDH QITNERGESSCDRLTDPHRAPS DTGSLSPPWDQDRRMMFPPPG QSYPDSALPPQRQDRFCSNSGR LSGPAELRSFNMPSLDKMDGS MPSEMESSRNDTKDDLGNLNV PDSSFP\AKKEATGPGFVPPPLA PVRGPLFPVDARGPFLRRGPPFP

SEQ ID		1	SEQ ID NO:		<u> </u>	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
20249	50617	Α	20365	160	2810	GLRVPLSPSSPRSRLRSGPLRLP
						GADSGSGPKAVCSPPFIVAPTG
						RGYCGDHESSFGAMEEPGATP
						QPYLGLLLEELRRVVAALPEG
						MRPDSNLYGFPWELVICAAVV
						GFFAVLFFLWRSFRSVRSRLYV
						GREKKLALMLSGLIEEKSKLLE
;						KFSLVQKEYEGYEVESSLKDAS
		İ				FEKEATEAQSLEATCEKLNRSN
						SELEDEILCLEKELKEEKSKHSE
						QDELMADISKRIQSLEDESKSL
	-					KSQVAEAKMTFQIFPMNEERLK
						IAIKDALNENSQLQESQKQLLQ
						EAEVWKEQVSELNKQKVTFED
						SKVHAEQVLNDKESHIKTLTER
						LLKMKDWAAMLGEDITDDDN
						LELEMNSESENGAYLDNPPKG
						ALKKLIHAAKLNASLKTLEGER
<u> </u>		1				NQIYIQLSEVDKTKEELTEHIKN
		1				LQTEQASLQSENTHFENENQKL
						QQKLKVMTELYQENEMKLHR
,						KLTVEENYRLEKEEKLSKVDEK
						ISHATEELETYRKRAKDLEEEL
						ERTIHSYQGQIISHEKKAHDNW
						LAARNAERNLNDLRKENAHNR
		ŀ				QKLTETELKFEL\LEEDPYALDV
						PNTAFGREHSPYGPSPLGWPSS
						ETRAFLSPPTLLEGPLTLSPLLP
						GGGGRGSRGPGNPLDHQITNER
						GESSCDRLTDPHRALSDTGFLS
						PPWDQDRRMMFPPPGQSYPDS
						ALPPQRQDRFCSNSGRLSGPAE
20250	50618	Α	20366	131	461	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20251	50619	A	20367		1761	MARGYGATVSLVLLGLGLALA VIVLAVVLSRHQAPCGPQAFAH AAVAADSKVCSDIGR\AILQQQ GSPVDASIAALVCTSVVNPQSM GLGGGVIFTIYNVTTGKVEVIN ARETVPASHAPSLLDQCAQALP LGTGAQWIGVPGELRGYAEAH RRHGRLPWAQLFQPTIALLRGG HVEAPVLSRFLHNSILRPSLQAS TLRQLFFNGTEPLRPQDPLPWP ALATTLETVATEGVEVFYTGRL GQMLVEDIAKEGSQLTLQDLA KFQPEVVDALEVPLGDYTLYSP PPPAGGAILSFILNVLRGFNFST ESMARPEGRVNVYHHLVETLK FARGQRWRLGDPRSHPKLQNA SRDLLGETLAQLIRQQIDGRGD HQLSHYSLAEAWGHGTGTSHV SVLGEDGSAVAATSTI\NTPFGA MVY\SPRTGIILNNELLDLCERC PRGSGTTPSTVSGDRVGGAPGR CWPPVPGEPSPSSMEPSILINKA QGSNLVIGCAGGELIISAVAQAI MSKLWLGFDLRAAIAAPILHVN
						NKGCVEYEPNFSQEVQRGLQD RGQNQTQRPFFLNVVQAVSQE GACVYAVSDLRKSGEAAGY

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
NO.	sequence	liou	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
20252	50620	I A	20368	341	2239	  LCASSCPFICPPIRPSVCPPAAPL
20202	30020	ľ.	20300	3		LLGCRAMARGYGATVSLVLLG
						LGLALAVIVLAVVLSRHQAPCG
						PQAFAHAAVAADSKVCSDIGR
						AILQQQGSPVDATIAALVCTSV
						VNPQSMGLGGGVIFTIYNVTTG
		ļ				AQWIGVPGELRGYAEAHRRHG
						RLPWAQLFQPTIALLRGGHVVA
						PVLSRFLHNSILRPSLQASTLRQ
						LFFNGTEPLRPQDPLPWPALAT
						TLETVATEGVEVFYTGRLGQM
						LVEDIAKEGSQLTLQDLAKFQP
						EVVDALEVPLGDYTLYSPPPPA
		ŀ				GGAILSFILNVLRGFNFSTESMA
						RPEGRVNVYHHLVETLKFAKG
						QRWRLGDPRSHPKLQNASRDL
					ł	LGETLAQLIRQQIDGRGDHQLS
						HYSLAEAWGHGTGTSHVSVLG
						EDGSAVAATSTIYTPFGAMVYS
						PRTGIILNNELLDLCERCPRGSG
						TTPSPV/SGDRVGGAPGRCWPP
						VPGERSPSSMVPSILINKAQGSK
						LVIGGAGGELIISAVAQAIMSKL
		ł				WLGFDLRAAIAAPILHVNSKGC
į						VEYEPNFS/QETVSCFVALAVL
						KLLASSDPATLAPKWLRL*EVQ
						RGLQD\RGQNPDPEGPFFLDVV
						PACV/SQEGACV*RPSRT*RKSG
			<u></u>			EAAGLLKTLLCPELEVLAPP
20253	50621	Α	20369	203	1518	RCYLQFLALLLTSTSARAAAAI
						AAAEEPAGSPSVMTRAGDHNR
						QRGCCGSLADYLSSAKFHL*LG
1						HSLSTWGDRMWHFAVSVFLVE
İ						LYGNSLLLTAVYGLVVAGSVL
						VLGAHGDWVDKNARLKVAQT
						SLVVQNVSVILCGIILMMVFLH
						KHELLTMYHGWVLTSCYILIITI
						ANIANLASTATAITIQRDWIVV
						VAGEDRSKLANMNATIRRIDQL
						TNILAPMAVGQIMTFGSPVIGC
						GFISGWNLVSMCVEYVLLWKV
						YQKTPALAVKAGLKEEETELK
						QLNLHKDTEPKPLEGTHLMGV
						KDSNIHELEHEQEPTCASQMAE
						PFRTFRDGWVSYYNQPVF/LGW
		-			1	HGSCFPLYDCPGL*LHHHRVRL
						HSGTEWFHPQYFDGSISYNWN
						NGNCSFYLATSKMWFGSDRSD
				<u> </u>		LRIGTAFLFDLVCDLCIHAWKP
20254	50622	Α	20370	1	497	
20255	50623	Α	20371	1	1575	

SEQ ID NO:	SEQ ID NO: of peptide sequence	t .	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20256	50624	A	20372	251	1351	SLKSTDGVKCGLSINIFHKRREN ILFGMGNPLLDISAVVDKDFLD KYSLKPNDQILAEDKHKELFDE LVKKFKVEYHAGGSTQNSIKV AQWMIQQPHKAATFFGCIGIDK FGEILKRKAAEAHVDAHYYEQ NEQPTGTCAACITGDNRSLIAN LAAANCYKKEKHLDLEKNWM LVEKARVCYIAGFFLHVSPESV LKVAHHASENNRIFTLNLSAPFI SQFYKESLMKVMPYVDILFGK*DRSWPTFAREQGFETKDIKEIA QKDTSPCPKMNSKGQRIRDLHP KGRDDT\IMATESEVTAFAGLGI KTQKEI\IDTNEAGDAFVGGFLS QL\VSDKPLTECIRAGHYAASIII RRTGCTFPEKPDFH
20257	50625	Α	20373	1	201	
20258	50626	A	20374	15	304	
20259	50627	A	20375	51	283	TWVKEAQNDGLCVCYIFIFNCQ MTGLDIEKD\QIIEMACLITDSD LNILAEVRDAHVIQSYFLKGTG NITIIFSSFFI
20260	50628	A	20376	2	520	QRRLRDWGRGCWSRVMLGGS LGSRLLRGVGGSHGRFGARGV REGGAAMAAGESMAQRMVW VDLEMTGLDIEKDQIIEMACLIT DSDLNILAE/GNSVHEDKKFLD KYMPQFMKHLHYRIIDVSTVKE LCRRWYPEEYEFAPKKAASHR ALDDISESIKELQFYRNNIFKKK
20261	50629	A	20377	2	206	EFGTRRHVPPCLINFFKDQGSP* LARLVSNSWPQVIHPPWPPTVL GLRGVGHCPWPRLGNSSWVW GGH

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20262	50630	A	20378	1	1401	MRSQKTSLYFSLEAKFYCSLRN
						ERIASCOIINDAGIQTKTLMKRS
						HVAYALKKPVEQGDSFELESK
						GFMAITIAVCIEQNLRHSSSLFIC
						MSSTGQSGTTNNASSLIPSYVFT
						PPHFNLLDKQGRTAGKSHCGE
						DRKEHNNACHQEEEAPDVNPL
						HGLQASGSPGKTQRIKSQALEG
						TRLYSHSEETRSGLIRLLRLRQR
						RLRDWGRGCWSRDMLGGSLG
						SR\LLRGVGGSHGRFGARGVRK
						RGAAMAA\GESMAQRMVWVD
				İ		LEITGLDI*KVQIIEMACLITDSD
						LIILAEGPNLII\KQPDELLDSMS\
						DWCK\EHHGKSGLTKAVKEST
						NYHCQQAE\YEFPALWYDQPD
						SLPGLCPTCQEIKFHE\DKKFP*
						QNTCPQFMKHLHYRII\DVSHC*
						KELCQTAGYPEEYEFA\PKKAA
		1				FSIGPLDD\ISENL\KKVPFFPEN
						NHLPRKKIRMKRKEGKL*KMG
						ENEKTVS
20263	50631	С	20379	180	248	
20264	50632	Α	20380	455	1133	GIEGWRGEADTLPAEGQDGGR
						RVSKGDIFSGGAETAQGWGVM
						GPFQAAPSKRAGREHRGGQRR
						VQHVGWYPRPGSDKGHGKGT
		1				QDSRVPQLPPHPPPIHLVSRHRG
						KLRHGFLRPMPEPRGLESGKTG
						NCFGELQH/RTSP*GRSGPREGG
						PRNIAQQGGCRGSACGRRSH\E
		!				ALRPRVWCGEGPQWTW\CAVC
		1				P\NRSAPGAGLAD\RQHPGESRA
2026	50.500		20001		110	WGETRLGEAGGAE
20265	50633	A	20381	214 3	412 257	DATVIITVNITTWEAALIEDEEAL
20266	50634	A	20382	ا	231	PATVIITVNTTWSAALISRESAL
						RSFPVKAAALSPSREGGSVTAT
						PNVRSMTSAVPIMRVSVQ\SKR *PEEQN*QETYPQTTSCR
20267	50635	A	20383	1	1569	TEEQN QETTYTISCK
20268	50636	В	20384	15	1744	
20208	120030	₽	120364	[13	1/44	

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20269	50637	Α	20385	2	7205	QASGSCWSSPFFSASTPSAAAT
						GVPGPTDMAAVLQQVLERTEL
						NKLPKSVQNKLEKFLADQQSEI
						DGLKGRHEKFKVESEQQYFEIE
			:			KRLSHSQERLVNETRECQSLRL
						ELEKLNNQLKALTEKNKELEIA
						QDRNIAIQSQFTRTKEELEAEKR
						DLIRTNERLSQELEYLTEDVKR
						LNEKLKESNTTKGELQLKLDEL
						QASDVSVKYREKRLEQEKELL
						HSQNTWLNTELKTKTDELLAL
						GREKGNEILELKCNL
20270	50638	Α	20386	120	7384	GWTRFPRVPPRESPAGGRARAT
						ERGDRGGGPGVWVLLVFAFLL
						RFYPVGRCHWGPWAPPTWAA
						VLQQVL\ERTELNKL\PKSVQNK
						LEKFLADQQ\SEIDGP*RGGHGE
						I*RWESEQQYFEIEKRLSHSQER
						LVNETRECQSLRLELEKLNNQL
						KA\LPEKNKELEIA\QDRNIA\IQ
						SQFTRTKEELEAEKRDLIRT\NE
						RLSQELEYLTEDVKRLNEKLKE
						SNTTKGELQLKLDELQASDVSV
						KYREKRLEQE
20271	50639	Α	20387	2	433	
20272	50640	A	20388	2	181	GFRRVSLEF*FPGVRWGMRKL
						GAPDLAPFSQM*SGGRAPCVAS
						*PPCRGLRLMRTGSRR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	I control of the cont	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20273	50641	Α	20389	l T	2422	MAGALAGLAAGLQVPRVAPSP
						DSDSDTDSEDPSLRRSAGGLLR
						SQVIHSGHFMVSSPHSDSLPRRR
					]	DQEGSVGPSDFGPRSIDPTLTRL
						FECLSLAYSGKLVSPKWKNFK
						GLKLLCRDKIRLNNAIWRAWYI
						QYVKRRKSPVCGFVTPLQGPEA
						DAHRKPEAVVLEGNYWKRRIE
	1					VVMREYHKWRIYYKKRVSGG
						GPGRPQSFPPAAAGYRPPRKIPG
				E		KGILTPELAPLGPSIQSRADSAT
						VWPQRLLAASLPRGRLRKPSRE
						DDLLAPKQAEGRWPPPEQWCK
						QLFSSVVPVLLGDPEEEPGGRQ
						LLDLNCFLSDISDTLFTMTQSGP
						SPLQLPPEDAYVGNADMIQPDL
						TPLQPSLDDFMDISDFFTNSRLP
						QPPMPSNFPEPPSFSPVVDSLFSS
						GTLGPEVPPASSAMTHLSGHSR
1	:					LQARNSCPGPLDSSAFLSSDFLL
						PEDPKPRLPPPPVPPPLLHYPPP
						AKQETVPEFPCTFLPPTPAPTPP
						RPPPGPATLAPSRPLLVPKAERL
						SPPAPSGSERRLSGDLSSMPGPG
				ļ		TLSVRVSPPQPILSRGRPDSNKA
ļ						LLGSFLGSPNSLLPETENRRITHI
				1		SAEQ\KRRFNIKLGFDTLHGLVS
						TLSAQPSLKVSKATTLQKTAEY
	,					ILML\QQERAGLQEEAQQLR\DE
						IEEL\NAAINLCQQQLPATGVPI\
						THQRFDQMRDMFD\DYVRTRT
						LHNW\KFWVVSSLKPMAGGLQ
20274	50642	A	20390	315	387	
20275	50643	Α	20391	570	1017	FCLHILQVPHTLAGHSPCAHG/R
						PPLPRF/HDRLPEEIQQQIVRETF
						HLVLKRDDNICNFLEGGSLIGG
1						SDYKLIYRHYATLYFVFCVDSS
						ESELGILDLIQLEQFKKDNEETG
						FGSGTRALEQALEKTKANIKW
		L				VKENKEVVLQWFTENSK
20276	50644	A	20392	1	1128	
20277	50645	В	20393	62	130	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
20278	50646	Α	20394	99	718	ENRGWQSMIQAILVVNNHGKP
						RLVRF\YQRFPEE\IQQQIVRETF
						HLVLKRDDNI\CNFLEGGS/L*L
			:			GGSDYK\LIYRHYATLYFVFCV
						DSSESELGI\LDLIQVFVGNSWIS
						CFENVV*IWI*TFH\MDK\VHYIL
			:	,		QEVVMGG\MVLETNMNEIVAQ
				:		IEAPNRLEKSEG\GLSAAP\ARA
						VSAVKNINLPEIPRNINIGDLNIK
						VPNLSQFV
20279	50647	Α	20395	3	370	SLHLAAILGETSTVEKL/YAAGA
						GLCVAERRGHTA/LHLACRVG
Ī						AHACARAL/LQPRPRRPREAPD
						TYLA/QGPDRTPDTNHTPVAL/Y
						PDSDLEKEEEESHTPLHVAVIH
						K/DVEMVRLLRDAGADLDK
20280	50648	Λ	20396	2	1097	SRQSPATGGRLRWAPDAALAM
						AVVACLGQAADSDKWGDSRLS
						SLGPDAAAPGGPGLGAELGPGL
						SWAPLVFGYVTEDGDT\FSAGT
		ŧ				EYMDLQNDLRQVSHEGWCRA
1						WGPGFPV*LPTACPLLLQTALH
1						LAAILGETSTVEKLYAAGAGLC
						VAERRGHTALHLACRVGAHAC
						ARALLQPRPRRPREAPDTYLAQ
						GPDRTPDTNHTPVALYPDSDLE
		İ				KEEEESEEDWKLQLEVENYEG
						HTPLHVAVIHKDVEMVRLLRD
						AGA\DLDKPEPTCGRSPLHLAV
						EGPRQADVLELL\LRAGANP\A
1						A\RMYGGRTP\LGSAMLRPNP\I
						LAR\LLRA\HGAPEPEGEDEKSG
		<u> </u>				P\CSSSSDSDSRLSPSAWSPLPPG
20281	50649	Α	20397	47	478	LRLRKLKAIMLAFGRGKTRNSE
						ESSDSQEVIWKDTWMFTALGH
						CGISMFGLWDRELTLTNYELFIS
}						SSPTPS*VPQRLCYVVSTFLMYL
						SLDHLYTGEMKAFLHVLDTQS
						VNSDFAFFVRLSFQYFYKLLCV
		<u> </u>				LWQGTVNKEFI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20282	50650	Α	20398	1	1992	METFVIGWQPAVTVWTIGVAP
	1					PPTLEITSQHEIWVAQHPNYIIS
						VKQESPAPMPWTAVALDSHRS
						MNTIVNCAREGSTLHAPYESLM
						PDDLRWNSFILKSSPSPHSRSME
						KLSSMKLVPGAKYIRDHCLVK
i						EALTIHKSPILLAAIDLLGNLSG
						FTDYQSYVIHGCLENNPTLERSI
						ALFNGISKWVQLMVLSKPTPQ
						QRAEVITKFINVAKGATAELNL
			]			WQHPKIIMLSQVVPGVFSQEHS
	į					LIRSLHANLHFTVCFPEDPLAM
	į					VAGMVQESTCPCEILQLPTTGQ
						LANWNEMTELVSSNGNYCNYR
						KPFPTCDGLKIPILGVHLKDWIA
						VHVNFPDWTEEIKVNIVKMHQ
						LSVTLSELVSLQNASHHLEPNM
		1				DLINLLTV/DGLQPADLMHAISL
						FRTSHRPQGLFL*IKGTSPAEHS
		•				DRSCQHFRV*KCSSWESWHGG/
						WLSLEE*MSSFLLRSF*KI/YCSR
						KNVACQTRAVCTNLDALLSSS
		ļ				QMSNNHHPFSLPAQDEVFEFPG
						VTAGHRDLDSRAITLVTGSSRK
						ISVRLQRATTSQATQTEPVWSE
		1				AGWGDSGSHTFPKMKSKFHDK
						AAKDKGFAKWENEKPRVHAG
						VDVVDRGTEFELDQDEGEETR
						QDGEPAEAVSVGAVAAPGMSE
						KELLQVDGRTPTRVLRMADLV
						GISEFRGEPAFV
20283	50651	Α	20400	1	417	
20284	50652	Α	20401	2	94	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20285	50653	Α	20402	190	2189	AFCPAGTRSAPCPRPRTMTMTL
						HTKASGMALLHQIQGNELEPLN
						RPQLKIPLERPLGEVYLDSSKPA
]						VYNYPEGAAYEFNAAAAANA
						QVYGQTGLPYGPGSEAAAFGS
				ŀ		NGLGGFPPLNSVSPSPLMLLHPP
						PQLSPFLQPHGQQVPYYLENEP
						SGYTVREAGPPAFYRPNSDNRR
						QGGRERLASTNDKGSMAMESA
						KETRYCAVCNDYASGYHYGV
						WSCEGCKAFFKRSIQELPTLC*I
						GWYWYQNRDIDQWNRTEPSEI
						IPHIYNHLIFDKRDKNKKWGKD
						SLFNK*CWENWLAICRKLKLD
						HFLTPYTKINSRWIKDLNVRTK
						TIKTLEENLGNTIQDIGMGKDF
						MSKTPKAMATKAKIDKWDLIK
						LKSFCSLLGYLKTENTAKETTIR
						LNRQPTEWEKIFAIYSSDKGLIS
						RIYKELKQIYKKK/GNPINKWV
		1				KDMNRHFSKEDIYAANRHMKK
						CSSSLAIREMQIKTTMRYHLTP
1						VRMVIIRKSGNDRCWRGCGEIG
						TLLHCWWDCKLVQPLWKTVW
		-		]		RFLRDLQLEIPFDPAIPLLGIYPK
						DYKSCCYKDTCTRMFIAALFTI
						AKTWNQPKYPTTIG*IKKMWHI
						YTMEYYAAIKKDEFI/SLVGTW
						MKLETIILSKLTQEQKTKHHMF
						SFITGSRTMRTHGCREGNITHW
						GLLRDSVNMVTQAH
20286	50654	Α	20403	2	907	RVPAMATSSFALPRILGAGARA
						PSRWLGFLGKATPRPARPSRRT
						LGSATALMIRESEDSTDFNDKIL
						NEPLKHSDFFNVKELFSVRSLF
						DARVHLGHKAGCRHRFMEPYI
						FGSRL\DHDIIDL\EQTATHLQLA
						LNFTAHMAYRKGIILFISRNRQF
		1				SYLIENMARDCGE\YAHTRYFM
						GGMLTNARLLYGPTVRLPDL\II
						FLHTLNNIFEPHVAVRDAAKM
						NIPTVGIVDTNCNPCLITYPVPG
						NDDSPLAVHLYCRLFQTAITRA
						KEKRQQVEALYRLQGQKEPGD
			L			QGPAHPPGADMSHSL
20287	50655	В	20404	70	1477	

SEQ ID	SEQ ID NO:		SEQ ID NO:	Nucleotide location of first		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
NO:	of peptide sequence	hod	in USSN 09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
20288	50656	A	20405	153	688	AHPLLRSSPEETSRSRKPRCATT
						NGFLRIPEKTTPWSPEVAIHRNR
		}				\ITPNKPPTVKSLEKVFC*LDKE
						AQKEKESQK*KGPVRMPTKT\L
						RITT\RKTP\CGE\GSKDRWDRF\
	·					QIEISQSRLH*LCTSPF*DLFKQI
						TFHQYLSPGV*GEKSPFADALS
						QISIHTIDDQFVKKKKKKKKW
20289	50657	A	20406	26	603	RPRGTLREYKVVGRCPAHSHK
1						CHTPPL\YR\MRNFAP\N\HVVA
				=		KS/RAFWYFVSQLKKDEESLPG
						EIVYCGQGV*ESSPCGSKNFGI
		1				WLRY\DSRSG\THN\MYRE*PGP
		1	İ			*PTAGGCSPSCYPRHGVARHAG
						PEAHFHFRFIERLEEI\AGQQNC
						RRPGCSKQFPRISRFKFP\LAPPG
:						SLRRQDKPRF\TTKRPNTFF
20290	50658	Α	20407	161	418	
20291	50659	Α	20408	1	2580	
20292	50660	Α	20409	3	264	
20293	50661	Α	20410	3	302	ALALLSPWVVDGTRHRGAGGG
						AHPGGSGRTGAHRGGGRLRHG
						GLQVPSPAPREGS*GPARNRAQ
						RRAVDRTQ*ECSSKEVYT*RLV
						PPPPLGHSHSSPE
20294	50662	A	20411	2	415	
20295	50663	С	20412	1	1230	
20296	50664	Α	20413	238	1765	VTAYHAATMNEVSVIKEGWLH
						KRGEYIKTWRPRYFLLKSDGSF
						IGYKERPEAPDQTLPPLNNFSV
						AECQL\MKTERPRP\NTFVIPCP
						QLTNIIRGTL/HTLVSPNQRE\EW
İ						NPPYQVVPQQPSRGGPKARTP WNNKCGSPK*LSHD*ENGSGGS
İ						ASHG/RKVTMNDSTYLNLLGK
ľ	1					GTFGKV\ILFRKKATGR*FA\MK
						\IFRKKVINAKD*VAHNIQPRAG
	1					VLPEHPGNPFLQLR*KYAFPDP
						TTRPVPL*MEYA\NGGELFF\HL
						S\RERV\FTEERA\RFYGAEIVSA
						SEVTWHSRGRGNTRDNQSWKT
						LMLEQKMAHIK\ITDFGL\CKRG
				1		ASSDGAT\MKTFCETP\EYLAHE
						\ILEDNDYGRAVDWWGLGVVM
						YEMMCGRLPFYNQDHERLFELI
				1		LMEEIRFPRTLSPEAKSLLAGLL
				1		*\KDPKQRLG\GGPSDAKEVME
						HRFFLSINWQDVVQKKLLPPFK
						PQVTSEVDTRYFDDEFTAQSITI
				1		TPPDRYDSLGLLEAGPADPASP
						QFSYSASIRE
L		<u> </u>	<u></u>	<u> </u>	<u> </u>	Kreisveige

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20297	50665	Α	20414	192	866	LSCRMRMLVLGQNPTLDSIDRS
						GVRSKCRIFNKLRESFHYNSAE
						LTISVNMSYNCCSGNFSSRSCG
						DYLRYPASSRGFSYPSNLVYST
						DLCSPSTCQLGSSLYRGCQEIC
						WEPTSCQTSYVEVQAPAKTSC
						YRPKNLPCSASPCKTTYS\GSLG
	1					FGSSSCRSLGYG\*GSCYSVGCG
						SSGFRSLGYGGCGFPSLGYGVG
						FCRPTYLASRSCQSSCYRPTCGS
20298	50666	Α	20415	312	1420	RSHSNAVPSVWNLTLPLDSGPH
20296	30000	Α.	20413	312	1420	QSPASTTGPCTNTERSSVPMSCS
	1					PPACPTQCVLCSCTEGQIYCGL
						TTCPEPGCPAPLPLPDSCCQACK
						DEASEQSDEEDSVQPLHVVRHP
						1 .
						QDPCSSDAGRKRGPGTPAPTGL
						SAPLSFIPRHFRPKGAGSTTVKI
						VLKEKHKKACVHGGKTYSHGE
						VWHPAFRAFGPLPCILCTCEDG
				•		RQDCQRVTCPTEYP\C\RHPREK
						WPGKCCKICPEDKADPGHSEIS
				1		STRCPKAPGRVLVHTSVSPSPD
						NLRRFALEHEASDLVEIYLWKL
	i					VKDEET*GSGEVKYLAQGHTA
		İ				RIFHLTQIKKVRKQDFPERGTA
						LPTARWPPRRSLERLPSPDPGAE
						GHGQSRQSDQDITKT
20299	50667	Α	20416	17	1008	KSGARSLQPQFLRAPEDGGPLS
			1			LPNAAMARGPKKHLKRVAA\P
		1				KHWMLDKLTGVFAPRPSTGPH
						KL\RECLPLIIF\LKDQTLSNALT
į		-				GD\EVKKI\CMQRFIKI\DGKVRT
						DITYP\AGFMDVI\SIDKTGREFP
						VL/YLIDTQGVRFCL*HRIYTLR
						EGQSTSLCKVRKILLWAPKGIP\
						HL\VTH\DAR\TIR\YPDP\LIQGE
						MNTHSRLI*KTGK\IT*FPSKFDH
						LVTCVMVT\GGA\NLGRNWVLI
						TQQRRSTLGSF*PLVHVK\DAN
						GNKLLATSDFSNIF/VVIGKGNK
						PW\ISL\PRGKGIPPHHLLEERDK
						RLAAKQSSWVKWGPWVTWSD
20300	50668	Α	20417	1	1377	
20301	50669	A	20418	1	2178	
20302	50670	A	20419	158	250	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20303	50671	Α	20420	ı	860	MDQTVKSLMASEEEYSTKEEK
		ł				YEKEMKLLEEKLKEAETRAEF
		i				VERSVAKLEKTTDDLEETLASA
		ŀ				KEETWRFIRPGTRPYWNSATCF
		ŀ				LLPGHKMAAEVHKLNPNVSEH
			ŀ			DQERWHDAFGNGGADNHSKL
		ĺ	,			QASTRRGGPLSQELLQSLAASL
						AAGGLTALSAEFTIGKGELMAH
ľ						DVLWGCAPDEYDDLISGTCSHL
						PKGKTHDVLTGSDDGRHSNTS
						LAWGAGVQFNPTESVAIDIAYE
		ĺ				GPGSGDWRTDGFIVGGLFCGV
						NMAVKIS\GVLKDGTGKPVQN*
			•			TIQLKAKRNSTTVVVNTLASEN
						PDEAGRYSMDVEYGQYSVILL
i						VEGFPPSHAGTITVYEDSQPGTL
					İ	NDFLCAMTES\DARPEALGRFG
				•		LYWWKEVARNAS\AVGQNTA
]						AAKKSASDASTSAREAATHAV
						DAGGFKNAQARTSSPDKAGSA
		l				GFKQGFPRGGNGINKRAT*RIK
		1				KVACR*SLMASEEEYSTKEEKY
		1				EKEMKLLEEKLKEAETRAEFVE
						RSVAKLEKTTDDLEETLASAKE
						ETWRFIRPGTRPYWNSATCFLL
		1				PGHKMAAEVHKLNPNVSEHDQ
		1				ERWHDAFGNGGADNHSKLQA
ľ		l				STRRGGPLSQELLQSLAASLAA
,		l				GGLTALSAEFTIGKGELMAHDV
ļ		1		•		LWGCAPDEYDDLISGTCSHLPK
		l				GKTHDVLTGSDDGRHSNTSLA
	ŀ	1				WGAGVQFNPTESVAIDIAYEGP
20304	50672	В	20421	34	800	WONGVQITALIESVALDIATEGI
20304	50673	A	20421	130	368	RQVITTLYIPYRRRHLSPPDPEQ
20303	30073	^	20422	130	300	PLNHKQNRKVNHGGTEFQVVV
						LPFPKMKINCYRRNQFFLFFFST
						VKVPVGWSMVPLQAATAGAR
[					1	L*NSDISPLKSQQTQDKFPSSKP
						PHIYIKK
20206	50674	1	20423	1	57	HHHKK
20306	130674	A	20423	1	31	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20307	50675	A	20424	85	874	GPRGLVSMAGGSVEVTPSTFTC YLRPPSASPPTPPSHPGLLPRSPA LCSLHTGARTLPLFGGGRCSSA VLPPRPGP/RAGSGSGQRERFHR FQPTYPYLQHEIDLPPTISLSDG EEPPP*QVPCHLQVR\DPKQQL\ ELNRE\SVGAPPNRTIFDSDLMD SARLGGPCPPSSNSG/ISAPTCYG \SGGRMEGPPP\TYSEVIGHLPG GPSFQH\Q\QSSGPPSLLEGTR\L HPHTHSRP*RSAAIWSKEKDKQ KGHPSLGSPGGPGLGLRR
20308	50676	A	20425	3	250	ESVSATEDLAPDAAQGEDNSEI KELL\RENEPLPACNGPAQAQPS PTTERAKSQEEVLPSSTTPSPGG ALSPSGQPSSSVDAA
20309	50677	A	20426	2	393	
20310	50678	Α	20427	1066	1824	
20311	50679	Α	20428	3	902	
20312	50680	Α	20429	1	2163	
20313	50681	Α	20430	1	1890	
20314	50682	A	20431	]	2882	MKQQLPVSYLQVIFNHWASRIS GARGAGWGPVRLGRLCADRAL DGESRPGGGGQSGVPASEASQ KAAHGRRLPVPARLLRCAHSA LEPGRGSASSPRNSTTSDNDQPP DYSFSKRCDSHSLGFSLVWESR NLTLPPTSKLLHFSAFHFFIYKC RAEGHKLYKDLKNFLSAVKVM HESSKRVSETLQFIYY\SEWYGH EELKAIVWNNDLLWEDYEEKL ADQAVRTMEIYVAQFSEIKERI AKRGRKLVDYDSARH

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20315	50683	A	20432		1791	MPGARTSSSGASENHRARGQG GGPQGVGRMAEGKAGGAAGL FAKQVQKKFSRAQEKVLQKLG KAVETKDERFEQSANNFYQQQ AEGHKLYKDLKNFLSAVKVMH ESSKRVSETLQEIYSSEWDGHE ELKAIVWNNDLLWEDYEEKL\ ADQAVRTMEIYVAQFSEIKE\RI AKRGRKLVDYDSARHHLEAVQ NAKKKDEAKTAKAEEEFNKAQ TVFEDLNQELLEELPILYNSRIG CYVTIFQNISNLRDVFYREMSK LNHNLYEVMSKLEKQHSNKVF VVKGLSSSSRRSLVISPPVRTAT VSSPLTSPTSPSTLSLKSESESVS ATEDLAPDAAQGEDNSEIKELL EEEEIEKEGSEASSSEEDEPLPA CNGPAQAQPSPTTERAK\SQEE VLPSSTTPSPGGALSPSGQLSSS ATEVVLRTRTASEGSE\RPKKT ASIQRTSAPPRRPPPPRATASPRP
						CSGNIPSSPTASGGGSA\TSPRAS LGTGTASPRTSLEI\*PNPE\PPEK PVRTPEAKENENIHNQNPEELC TSPTLMTSQVASEPGEAKKME DKEKDNKLISADSSEGQDQLQV SMVPENNNLTAPEPQEEVSTSE

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nino acid   *=Stop codon, /=possible nucleotide
the same of the sa
deletion, \=possible nucleotide insertion)
QLQSKMKLFWLLFTIGFCWAQ
YSSNTQ/SKGRTSIVHLFEWRW
VDIALECERYLAPKGFGGVQVS
PPNENVAIYNPFRPWWERYQP\
VSYKLCTRSG\NEDEFRNMVTR
CNNV\GVRI\YVDAVINHMCG\N
AVSAGTSSTCGSYFNPWKLRTF
PAIP\YPGWDFNDGKC\KTGSGD
IENYNDATQVRDC\RLTGLLDL
ALEKDY\VRSKIA\EYMNHL\IDI
GVAGFRLDASKHM\WPGDIKAI
L\DKLHN\LNSNWFPAGSKPFI\F
QEVIDLRVGEPNLKASDYFGNG
RVTEFKYGAKLGTVIR\KWDGE
KMSYLKKWG\EGWGF\MPSDR
ALVFVDNHDNQRG\HGAGGASI
LTFW\DARLYKMA\VGFMLAHP
YGFTRVM\SSYR/WAKHSFKKG
NDVNDW\VGPTQINNG\VIKEV
TINPRH*LVGND\WV\CEH\RWR
\QIRNM\VNFRNGSGMAQPFLQ
NWYDNGSNQVAF\GRGNR\GFI
VFNK*WPVIFI*LCKPGLPAGSY
CDVISGDKVIGNCSAIKIPVSVD
GKAHFSMSNSAEDPFLAIHAES
KIVKFKIKCKSAKQKKKKKKG
RPF
CASGSSGWRPVLWAGAFTMAS
AELDYTIEIPDQPCWSQKNSPSP
GGKEAETRQPVVILLGWGG\SK
DKNLVKYSAIYHKRKLLELLFD
YEIEKEPPAFSLSSATVASCCTA
TCWSSCRPVASAACV
CASGSSGWRPVLWAGAFTMAS
AELDYTIEIPDQPCWSQKNSPSP
GGKEAETRQPVVILLGWGGCK
DKNLAKYSAIYHKRGCIVIRYT
APWHMVFFSESLGIPSLRVLAQ
KLLELLFDYEIEKEPLLFHVFSN
G/VRHAVPLTCWELLQTRRFCR
LRVVGTIFDSAPGDSNLVGGSA
GLGSHPGARG

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20320	50688	A	20437	78	1237	CWSHSDTYWPRTPRPPPPGRPG PVVHLAQPVLHPEAGIPEFWES GSPGQTRGAALVLTVHICVSEN SPSPGGKEAETRQPVVILLGWG GCKDKNLAKYSAIYHKRNTAT LQPPLTYMRGWGLFLTGAPAD QYLIQIKRISQWLRGCIVIRYTA PWHMVFFSESLGIPSLRVLAQK LLELLFDYEIEKEPLLFHVFSNG GVMLYRYVLELLQTRRFCRLR VVGTIFDSAPG*QQPG*GL\SGP GQPSWERRARPMLRLLLLVAF ALGGSSCFHVLLAPITALFHTHF YGQAYRTRALAWPELYLYSRA DEVVLARDIERMVEARLARRV LARSVDFVSSAHVSHLRDYPTY YTSLLCRTSMRKLRPLLKAIAPS
20221	50690	_	20429	2	427	HLCSQKINALKPSP
20321	50689	A	20438	3	427	
20322	50690	В	20439	101	1099	
20323	50691	Α	20440	733	2228	

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SEQ ID	SEQ ID NO:	ı	š	•		Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20324	50692	Α	20441	582	2595	PPPPVRRRDPGPAVRPAGAMA
						KKSAENGIYSVSGDEKKGPLIA
						PGPDGAPAKGDGPVGLGTPGG
						RLAVPPR\ETWTRQMDFIMSCV
		ŀ				GFAVGLG\NVWRFPYLCYKNG
				•		GGVFLIPYVLIALVGGIPIFFLEIS
						LGQFMKAGSINVWNICPLFKGL
						GYASMVIVFYCNTYYIMVLAW
						GFYYLVKSF\TSTLPWATCGNY/
						TWKHSPTCVEIF\RHEDC\ANAS
						LANLTCDQLANR\RSP\VIEFWE
						NKVLKLSGGLE\VPGALNWE\V
						TLGLLACWVLVYFCVWKGVK
				:		STGK\IVYFTA/THYPYVVLVVL
						LVRGVLLPGALDGIIYYLKPDW
						SKLGSPQVWIDAGTQIFFSYAIG
						LGALTVLGSYNRFNNNCYKDA
						IILALINSGTSFFAGFVVFSILGF
						MAAEQGVHISKVAESGPGLAFI
		ļ.				AYPRAVTLMPVAPI\WAALFFF
						MLLLLGLDSQFVGVEGFITGLL
						DLLPASYYFRFQREISVALCCA
						LCFVIDLFMG\TDGGMYVFQLF
						DYYSASGTTLLWQA\FWECVV
	1					VAWVYGADRFMDDIACMIGY
						RPCPWMKR\CWSFFTPLVC\MGI
						FILNVVYYEPLVYKQN/TYVYP
						WWGE\AMG\WPFALS\SMLFVP
		l				L\HLLGLPSLRAKGHPWLKRW
			]			QH\LT\QPIWGLHPL/EELPRFRD
						ADVQGTLTHP*TPVSESSKVV\V
20225	50602	_	20442	1	363	MGKRGLILLIEACIAIAQPLQRW
20325	50693	A	20442	1	303	FVVPPRAGMPVTYD/HSNSAAG
1						SPPV*NARLCSPGEKVLCRSQT
		1				•
						GNFPHTPQEQAVPSAWNALPV
						QP*SH/GPSPMLVPCRCVCPMET
20226	50604	_	20442	7.47	1200	RVQGSASPGGQG
20326	50694	A	20443	747	1389	LGAVQLCPPPCRPCHRAALPGG
1		1	1			LFPPPRPPHSPQAGPLHPQWTQ*
						PGSVGSPTQLSPEASRPQWPPT
						A\GSPRAGVHPGHGQA*SADISS
						AKQWTQVGPAGQGGSKFGKT
						KVPWQSGPLGWDPGARAGQG
						VG*ETRGPSLQLAWSDGEGV/P
						GWRGSYRGHRRGP*PPAPWPW
						SGGHPGSPMEWPCPGTELQMR
		1				NRAGSRDLHSLGRAPPAATAPS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20327	50695	Α	20444	90	1181	RAEPHLQVPGVPGSAQKPQGT
ļ						QMAPVSPRRGPQRGWHTLGAQ
						RLLPQAALSSPAFPQHPGGRWA
						PRRRGQSSQGHSPESLPSERAQP
			:			HWNHPQLLSPCTPLSCP*GGPR
						QA/EQAGDSGSGTWVGLGAAP
						SERGAPMSARHWLRAPGRRPIF
		i				LLSALW*TPAWGGGPAASRSP
	-				İ	GGPSSRVVGCPLCPLAPGPVPS
						VE*RPMSTKDLPPPSPR*KP/PA
						GPAPV*/SPKAVGIPAWERPPTR
						PETEGAGGSCAERLLAPVEMAP
						VSRGQRRDPEPGSRHG/DPPTDT
						AALGGPRPPQHLPRTRQHWEI*
			1			GPGAPPTDTAALGDLRPQSPGL
						LMELG\PISRDPHGRGPWQGPV
						GKLWPGGQGSCPQSPPV
20328	50696	Α	20445	2	957	IPSRKDHEKAEFEVHEVYAVDV
		1				LVSSGEGKVRRVPELAKRGD*E
		1				CSPDQMLLKLLFQAKDAGQRT
ļ		1				TIYKRDPSKQYGLKMKTSRAFF
		l				SEVERRFDAMPFTLRY*AMIVL
						GTSLTHRPYTQEHFFLSLPA*TC
			}			SVHVLTPSLSLPISFSLKVM*KS
		ł	ŀ			NPKHDFCLRVGAILKYEQNGK
	İ	ļ				T*WGVKDTSEYHLPCQ\AFEDE
İ		ļ				KKARMGVVECAKHELLQPFNV
		ŀ				LYEKEGEFQKELHFGFPDYKISF
	1	ŀ				L*VQKS*V*QRNFLSKHIFIFAIG
						EFVAQFKFTVLLMPNGPMRITS
				-		GPFEPDLYKSEMEVQDAELKA
						LLQSSASRKT
20329	50697	Α	20446	85	266	

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SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20330	50698	A	20447	207	1759	LSQRALRLSPRARSFSLSPACPL PCLLALSLALSSRIEGTLTTACG WGRETEAAAAQGKRGCSGGSR KMSGEDEQQEQTIVD/DSLVVT KYKMGGDIANRVLRSLVEASSS GVSVLSLCEKGDAMIMEETGKI FKKE\KEMKKGIAF\PTSISGTIN *PI/CHFSPFERSD\QDYIPPRKGD LVK\IDLGVPCWMGFIANVASH/ SFVVDVAQGTQVTGRK\ADVIK AAHLCAEAA\LRLVKPGNQNT QVT\EAWNKVAHSFNCTPI\EG MLSHSLKQHVIDGEKP*FQNPT DKQK\RAHEKADFEV\HEVYAV DVLVKPQERARPKDAGQRTTIY KRDPSKQYGLKMKTSRAFFSE VERRFDAMPFTLRAFEDEKKA RMGV\VECA\KHELLQPF\NVLY EKEGEFVAQ\FKFTVLLMP\NGP MRITS\GPFEP\DFYK\SEMEV\Q DAELKALLQ\SSAKSEKPQKKK KK\KASKTAGEWPLVAFSAVLE TFRRKWKLGGLEVGPISPACCS
20221	50.000	<u> </u>	20440	1	2025	CLHPPSHQTPRLCEVQFFFST
20331	50699	A	20448	1084	3825 2429	ARERSLRGSPAREREKERKCTA LSE*GEQMERKTNS*TKQTE*N RNEIGPRGAGPPGDAGSGRQPA LRPHVHKQTEKTTRVRTVLNE KQLHTLRTCYAANPRPDALMK EQLVEMTGLSPRVIRVWFQNK RCKDKKKSILMKQLQQQQHSD K/SGEQPLGRRLESGGGCASAA VSACPFLGEPWEIQGELGAPDG VCLYLHSKASLGFEASYFVWD RGFSLSQWQPRVAGSGRCAEP KAAWQCAAPASRAGGLCTPSA AAERSPVEKEGSSDAIECAPPR KAMQTLNHSRPGAPVSKVAEK ARPNSSTYCLLLAPQSLQGLTG TPLVAGSPIRHENAVQGSAVEV QTYQPPWKALSEFALQSDLDQP AFQQLGLLSAGGRSPRDTFRLG ARGPGAPARVPLHLPLPGVSAA P\VSFSESGSLGNSSGSDVTSLSS QLPDTPNSMVPSPVET
20333	50701	Α	20450	133	276	
20334	50702	A	20451	2	205	WDTSS*SLRTPRSRQVSTRPRR KKEKLPSKGMFSSVSLPQVLTQ RHQRPTCSWSLCWQPQHLILLR S

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20335	50703	A	20452	182	411	GRAWPAPPTSVTPRLRSPAESP RWPSAEERSSLP/VRGRRPGYG* VSCCGPRAPPGVNGDALRSARP AQLPA*RPQRH
20336	50704	A	20453	786	1208	GVHRGTHGRQKWRPCAPRAQ HSWGPGGRLPAGCRG*AGSCA/ PSLGL*CGAGPQGWGRRDP*TR RRTPRRPPRGPRPRVWMAPPAR SHPWRAAPRSAGAAPRPLGLG GRSGRETRPAPGGELQRPAAAG PPLMSCGWRVVLA
20337	50705	A	20454	385	727	KAMSPVFPRHSKWDLDLSRPLF TSAALLSACKILKLKVDKNKM VATSGVKKSYI*STV*TTREDW TAGRQRTWRCIYSTTEEKEDSG *SPSKGNGEGRGDAT*TTER*RS DTGL
20338	50706	A	20455	568	3382	HPGWVARGPGAGEGHAAGCG PARCQAGGPLQDGGCGGQGHR FGGGHVPEIKGGGFPHAGACAL EERQPR*QP*GVGKRGGGLFHA SSSPSSGGSRTEPAEG*GGAVCR AGDLGHIDDAGDHGDSGK\KE RVEEEGGKPKHVLSTSGVQSD AREPGEESQKADVLEGTAERLP PIRASGLGADPAQAVVSPGQGD GVPGPAQAFPGHLPLPTKVEAK APETPSENLRTGLELAPAPGRV NVVSPSLEVAPGAGQGA
20339	50707	A	20456	762	1245	LRQYSVKMRIVPTILLNFGADP DLRDIRYNTVLHYAVCGQSL\S LVEKL\LEYEADLEAKNKDGYT PLLVARY*QLIPKLVKFLLE\KG ADVNASDNYQRTALILAVSGEP PCLVKLLLQQGGEICNEG\MVD SQLRNMFISMVLLHRYPQFTAS HGK\QKHAK
20340	50708	A	20457	346	608	PKRRGAPLPVIPQSAS*KPKLPM GRPPPPVPSVSPGPLPGSLAIAP HSLEPHLVQSSCLVVPPKTKQS KRKPGRGAQAPTLPNATEFSVC RNDPKEGGRRSP*FPNRLLEPE APPRPSAGESPRPAAA/SPLPCSE EK/PTPQLCARERLAPSAAPPPP PRCGPAPLLEPSAWTPELARRA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
20341	50709	Α	20458	340	889	SAPFSSGERTPGCPEASPFPGGG
20311	30707	` `	20150	3.0		GGNARPSGPAPHPPSSAS*KPKL
						PMGRPPPPVPSVSPGPLPGSLAI
						APHSPEPQLVQSSCLVVPPKTK
						QSKRKPGRGAQAPTLPNATEFS
						VCRNDPKEGGRRSP*FPNRLLE
						PEAPPRPSAGESPRPAAA/SPLPC
i						SEEK/PTPQLCARERLAPSAAPP
						PPPRCGPAPLLEPSA\EAEEGGW
						GAGPLGRALPPPPPGKGLASGQ
						PGVRSPDEKGADQRQPSGPRLC
		1				SAPAPPRRS/PPRRRRPSNFVPED
						QPHYVPGSASAGPRRSPRPPRR
						ELGAPSHLP
20342	50710	Α	20459	148	397	GHKELVTGTEAIELAHAYETCH
			,			AVPRTIAKLVPKTH\LMSESEW
						RNLGVQQE/RQGWVH\YMIHEP
						E\P*HFGCSRRPLTQGTQRK
20343	50711	Α	20460	139	272	VLKPGGSYSFGSPSCNSSH*LIG
ŀ						FQKWKDVPCEDKFSFVCKFKN
20344	50712	Α	20461	440	1167	ARDSLQLSMAQTSSYFMLISCL
1		1				MFLSQSQGQEAQTELPQAR\ISC
		1				PEGTNAYRSYCYYF*WKTRET
}						W\VDADLYCQNM\NSGKLGVC
						GSPRAEGCLFWPQLI*GGVGT*
						WTFNVLDLGLQ*PPKRTRRW\H
						WEQWGPLVSLQSSWGHWEPPS
		ŀ				RCLNPG\Y\CVSLDLQSTRISRN
						WKDVPFVKDK\FSF\VCKVQKT
		ļ				RGQAGKIQCLELTPAITNGVQK
						LNRTISPTQLNLVTLFSAEFALLI
20015		<u> </u>	20162		200	FNSFT
20345	50713	Α	20462	3	322	DI VELVOVA CNUDDI VI LIZVO
20346	50714	Α	20463	I	420	DLYELVQYAGNIIPRLYLLITVG
	1					VVYVKSFPQSRKDILKDLVEMC RGVQHPLRGLFLRNYLLQCTR
						NILPDEGEPTDEETTGDISDSMD
						FVLLNFAEMNKLWVRMQHQG
						HSRDREKRERERQVSW\RCKCG TLQ\QIVLTG
20247	50715	_	20464	1	1446	TEQ/QIVETO
20347	50715	A	ZU404	1	1440	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
ļ				sequence		
20348	50716	A	20465	3	2564	LGAGAACCRLWGVAMPTTQQ
20340	30710	<b> </b>	20103	ľ		SPQDEQEKLLDEAIQAVKVQSF
		ĺ				QMKR\CLD\KNKRIGVRHKTML
		1				RNMLG\ELRTSMLSPKSYYELY
	1					MAISDELHYLEVYLTDEFAKGR
						KVADLY\ELVQYAG\NIYPQGF
						YLLVHSLSCNNVKSFPQSR\RDI
						LKDLVEMCRGVQH\PLRGLFLR
	1					NYLLQCTRNILPDEGEPTDEETT
						GDIRDSMDFVLLNFAEMNKLW
						VRMQHQGHSRDREKRERERQE
						LRILVGTNLERLSQVE/WCKCG
						TL\RQIVLTGILEQVVNCRDALA
1	-					QEYLMECIIQVFPDEFHLQTLNP
						FLRACAELHQNVNVKNIIIALID
						RLALFAHREDGPG\IPADIKLFDI
		ĺ			i	FSQQV\ATVIQSR\QDMPSE\DV
						VSLQVSL\INL\AMKCYPDRV\D
						YV\DKVLGNNSGRIFNKLNLEHI
		1				ATSSAVSKGNFTRLLGKLPVGH
		1				FTNNILT\VL\KLKHFHPLFGYFD
		ļ		:		YESRK\SMSCYVLS\NVL\DYNT
						EIVSQDQVDSIMNLVSTLIQ\DQ
						P\DQPVEDPDPEDF\ADEPEALV
						ARFIHLLRSEDPDQQYLILNTAR
						KHFGAGGNQRIRFTLPPLVFAA
						YQLAFRYKEI\SKVDDKWEKKC
		Ì				QKIFSFCPTKTI\SALIKAEL/AQE
						LPLRTFSFKGGT*LA/AGELGFG
						NHG\TVAY\EFMSQGIFSVWKD/
						EKSGDFPKAQLGWPFTFGSIGA
						LFERVEVLSVEEDSWNPLR\TQ
20349	50717	С	20466	23	385	
20350	50718	С	20467	16	435	
20351	50719	Α	20468	1	436	
20352	50720	A	20469	1	747	
20353	50721	Α	20470	1	864	

SEQ ID NO:	SEQ ID NO: of peptide sequence	l .	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20354	50722	A	20471	2	2138	VGRAAAAAAAVAVPLAGGQE GSPGGGRRGSRGTTMVKKRKG RVVIDSDTEDSGSDENLDQELL
	-					SLAKRKRSDSEEKEPPVSQPAA\ SSDSETSDSDDEWTFGSHKHKK
						KGQARTLEQKGTMKKQANKT ASSGSSD/RTDSSAESSAP\EEGE
						VSDSDSNSSSSSSDSDSSSEDEE
						FHDGYGEDLMGDEEDRARLEQ
1						MTEKEREQELFNRIEKREVLKR
						RFEIKKKLKTANKKEKKEKKK*
						QEEEQEKKKLTQIQESQVTSHN
1						KERRSKRDEKLDKKSQAMEEL
				İ		KAEREKRKNRTAELLAKKQPL
						KTSEVYSDDEEEEEDDKSSEKS
		=				DRSSRTSSSDEEEEKEEIPPKSQP
						VSLPEELNRVRLSRHKLERWCH
						MF\FFAKTVTGCFVRIGIGNHNS
						KPVYRVAEITGVVETAKVYQL
						GGTRTNKGLQLRHGNDQRVFR
						LEFVSNQEFTESEFMKWKEAM
						FSAGMQLPTLDEINKKELSIKEA
						LNYKFNDQDIEEIVKEKERFRK
						APPNYAMKKTQLLKEKAMAE
1						DLG\D\QDKAKQIQDQLNELEE
1						RAEALDRQRTKNISAISYINQRN
						REWNIVESEKALVAESHNMKN
1						QQMDPFTRRQCKPTIVSNSRDP
						AVQAAILAQLNAKYGSGVLPD
						APKEMSKASVGQGKDKDLNSK
		ŀ				SASDLSE\DLFKVHDFDVKIDLQ
						VPSSESKALAITSKAPPAKDGAP
20355	50723	A	20472	438	709	FLSQSWIFKQIRASRTLFLVAFS
1						SNTFTSCRSFCSSSEAMGAGSG
	,					RSPELQPRRRRNRKPRAEPSNR
	İ					AQTPTRTS/LRRCSLRPHRPCAR
						SR
20356	50724	Α	20473	2	1315	ILQKRIGGFLDALVRNSQPARYI
		1				NDYQLSNYGLDMLGPSLDPAP
1						MASEELQKDLEEVKVLLEKAT
						RKTEKSKIETEIKNKLQQKSQK
		ĺ				KAELLDNEKPAAVVAPITTGYT
		1				KGGGGEERGGEAPTSNEAQAT
						EVRGGAGGKDKDDGDRER\P
						RKTRGGRARTE*NNRHEQ*IQI
						KKRRETEKERRNEKGRRRHKT
						ATQKRRREERRARRVGEGTRK
						REQPAHEQEHATTHTRPPAP/R
20255	50705		20474	42	1.60	RTRKTQTHNRKERKKEE
20357	50725	С	20474	43	162	

SEQ ID			SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20358	50726	Α	20475	2	363	YEPLDSNKRIGTSSSSYVLCLCS
						SLAIFSNFMGNLLSFCGVLKLK
İ						SLACIH*RFFFN*A**VFIHISSRE
						Q*FINVCSSCPAFFNPERRNELY
						ASLLSFVPSALVPIFEDNVSYLF
						VKSF
20359	50727	Α	20476	505	2294	VYSDAKSIYCLLFAHSQRPPCV
						FNENDPQNSGTVHTVFLVGLTD
						HRHARAKIESWAGEAAPFTLV
				-		AASADKLFGDLHISKYCGRIAV
						SVFQFQQYCGLCDSTS/SLQQY
		1				CGLCGST*HSGSTVASVAGLHI
						VAVPWPLWQDFTLWQYRGLC/
						AQDFTLWQYHGLCGRTSRVAV
						PWPLWQYFTLWQYCGLCGRTS
						HC/GQYRGLCGRTSHCGSTVAS
						VAGLHVWQYCGLCGSTSRC/E
						QYHGLCGRTS/RLWQYHGLCG
						RTSYC/EQYRGLCGSTSRCG/MY
						RGLCGRTSH/WWQYCGLCSSTS
		1				YCSRTVA\LWQYFIL\CSTSY/W
						QQYHGLCGNTSYCGIL\QLQQY
						SGLCSSTSYWGSTVASVAVLHI
						AAVLWPLRQYFILQQNRGFCG
						NTSYCG/SYFILWQYHGLCGST
						SYCG/SYFILQQYHGLCGSTSYC
						/EQYRGLCGSTSYC/TQYCGLCG
1						SSSYC/AQNCGFCGNTSYCGSTS
						/VLQQYHGLCGSTSYCGRTV/V
						FCGSTSYWGSTSYCGSTVASLA
						VLHIVAVLCPL*QYFTTNKTKN
						KTKPGSCHLTYFGGAAGGPSL*
						APGGVAGTVGSASRSASWGLP
ŀ						EWICCLRDPKSLSFPYWGFEVW
20360	50728	Α	20477	2	554	LSSTGGLHAGRLCCSPCLLVIPL
						KSSQHIF\RVL\NTNIRLGRRKNS
						PLPITAHLRVWARRYA\HVVLR
						KADIDLTK\RAGEL\TEDE\VER
						VITH/ILQNPRQYKDPKTGFFEQ
						DKKDVK\DGKYSQVPSQLVWD
						NKLP*KTWEPTERRFRAH*/RGL
						RHFLGAFRVRGQAHQEPLGRR
						GRHRWVCPRRK
	1	Щ_		L	1	L=

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20361	50729	A	20478	50	774	LFGVVYTGFCLGVWTAGVGGV PGPNFPRPILTLVRGP*GKGVG WVAGPGQLRPTGPWSI*VSHPT PPGPLSISYRAQAAGGQGPAP\I THPQGRQEP*RQLPPGGPLA/RH PPKAPDPARPHP/RQHRGPS/ST APKGRCWQHKQEPSGHPAGHT PGRPG/PPQAPPRPSPWVGPGPQ TPTPCP*DEPHRPPGLQPWRPFS PLAMAPMGRLQALPGLDQRGQ TVSPLRLLGALESLAWLPSAFIA
20362	50730	C	20479	156	908	
20363	50731	В	20480	1	804	
20364	50732	С	20481	278	551	
20365	50733	A	20482	169	1144	INYSLEKHVGALGRVLFSL*RA GCPGMGSTRERGLYLGKHRGS GGIW*ALAGP*KSRGDSVSLTQ GHTHVCSRSPR*ADSPPG/SHLS PVPHSVEVAGHVLVPATRAAV PCSASAGA*QSTYRTGVHQGNP TV*TK/PSRRPSGGVAK*FLPSA VRGEPGAKPLVDDLLPGWSLA THGQPPLVAAPGSGLWGRPAD A*GCETAGGSPCPRSTSRPSGPS GVQGCPLG*AGSGASASRSEPP GSTSCCPRAP\T*PAAPCVPDWP AGDQWRSHGYLPPSREL*G/W MPPSRPATLPQLAFARQRQGNR FDAAFESSGEDFHQMPRVGRM
20366	50734	С	20483	31	90	
20367	50735	В	20484	48	197	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20368	50736	A	20485	5447	5642	PRDMESCYNPGLDGIIEYDDFK LNSSIVEPKEPAPETADGPYLVI VEQPKQRGFRFRYGCEGPSHGG
					į	LPGASSEKGRKTYPTVKICNYE
	•					GPAKIEVDLVTHSDPPRAH\AHS
		1				LVGKQCSKLGICAVSVGPKNM
		į				TAQFNNLGVLHVTKKNMMGT
						MIQKLQRQRLRSRPQGLTEAEQ
			1			RELEQEAKELKKVMDLSIVRLR
l						FSAFLRASDGSFSLPLKPVTSQP
						IHDSKSPGASNLKISRMDKTAG
1						SVRGGDEVYLLCDKVQKDDIE
						VRFYEDDENGWQAFGDFSPTD
						VHKQYAIVFRTPPYHKMKIERP
						VTVFLQLKRKRGGDVSDSKQF
						TYYPLVEDKEEVQRKRRKALP
						TFSQPFGGGSHMGGGSGGAAG
1						GYGGAGGGGSLGFFPSSLAYSP YQSGAGPMRCYPGGGGGAQM
						AATVPSRDSGEEAAEPSAPSRT
						PQCEPQAPE\MLQRAREYNARL
						FGLAQRSARALLDYGVTADRR
		1				ALLAGQRHLLTAQDENGDTPL
						HLAIIHGQTSVIEQIVYVIHHAQ
1						DLGVVNLTNHLHQTPLHLAVIT
		İ				GQTSVVSFLLRVGADPALLDRH
						GDSAMHLALRAGAGAPELLRA
İ						LLQSGAPAVPQLLHMPDFEGLY
	1	1				PVHLAVRARSPECLDLLVDSGA
						EVEATERQGGRTALHLATEME
						ELGLVTHLVTKLRANVNARTF
				į.		AGNTPLHLAAGLGYPTLTRLLL
20369	50737	Α	20486	3	346	PQTITDTELWVTLTVEVWDIST/
						LFPGN*SHA/TLPSH*NLITLTPL
1						NANI/AIPQHALKGLKPVITRLL
						QHGLLKPINSPYNSPILPVQKLD
1						KSYRLVQNLRLINKIVSPIHPVV
						PN
20370	50738	Α	20487	1	808	YGTSLLLPWQLITCPLPSH*NLI
						TLTPLNANI/AIPQHALKGLKPVI
						TRLLQHGLLKPINSPYNSPILPV
						QKLDKSYRLVQNLRLINKIVSPI
						HPVVPNPPAFTSQITQAVSQAL
						GIQWNLHIPYHPQSSGKVERTN
						GLLKVHLTKLSLQLKKDWTVL
						LPLALLRIRACPRDATGYSRFEL
						LYGRTFLLGPNLIPDTSPLGDYL
						PVLQQARQAANLLLPTPDPQPH
						EDTLAGRSVLVKNLTPQTLQPR
	<u> </u>				L	WTGPHFIIYSTPTAVCLQDPPH

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,217	codon for peptide sequence	or peptide sequence	deterion, -possible nucleotide insertion)
20371	50739	Α	20488	2	513	CCPIASEAQ*TITDAELWVTLTV
						EAQSLLRVFLSYLPTLTHKYGT
						SLLLPWQPITCPLPSR*NLITLTP
						LN/GQYPIPQHALKGLKPVITRL
						LQHGLLKPINSPYHSPILPVLKP
						DKAYKLVQNLRLIYQIVLPIHP
						VVPNPYTLLSSVPPSTTHYSVL
						DLKHAFFTIPLQP
20372	50740	A	20489	1	1893	
20373	50741	C	20490	1	1806	MESCADO LO DAVIA LA TRADACIONA
20374	50742	A	20491	1	645	MTSGPQINQPKKHLTNFKSEPS
						DL/FPSSPGCSLPAED*RCPIALE
						AP*TITQA\RVTLIVEGKSVPLV
						NTEATQSTLPSFQGPVSLASITV
						VGIDGQASKPLKTPLLWCQLG
						QHSFMHSFLVIPTCPVPLLGRDI
						LTKLSASLTIPGLQPHLIATLLP NPKPPSCLPLLSPHLNPQVWDIS
						TSLATDHMPITIRLKPNHPYPAQ
						`
20375	50743	A	20492	113	328	RQYPIPQHALKD LGSGAWGGGDLPWEINPLSSCS
20373	30743	^	20432	113	328	LLREKDLLTTSGPQT\TSPRNISP
						ILNRDPTVQLTWQPLPEPLELW
						PKAL
20376	50744	С	20493	183	254	TATE
20377	50745	Ā	20494	1	511	MWKGPKGLDMYGKSSVSPKTS
						DILGRDTLLLALKVQTVVLQTA
						CGEGHVAGNCGRPLETEGSLQ
						LTATKKLRDSVLQPKSPEFCQQ
						FTRAWNRTQVPDETEAPAGTY
						AAQSGDLPWEINPLSSCSLLHE
						KDPPTASGPQT\TSPRNISPILNR
		}				VSEVSDHAGTPALVLHP
20378	50746	Α	20495	1	1108	MWESLELPGNIFNGFDQNADN
						DMDDEIQAEVISDGDEEFVGN
		1				WNKPRDLVPCILATLALTKRGQ
						DTAQTMASEGARPKHWQLPGG
						VGPVGAQKSRIEVWEPLPIFRR
						MYGKACMSRQKFAAGAGFSW
						YVPVAVVGAKVHDVNLHMLS
						FPSKWKLHTCMKFGAVTQIVTS
						LGRSSCSLLLEKDPPMVLRPTSP
						RNISPISNLTKETRFIRGPKTPAP
						VTDWEGSLPLVFNHCRDASLII
						HPGFRGVRPRRDACLSPSPLAN
						LINLTFKVYNNRKKLQFLAFTV
						RQTSAMSPAHKNFQTPEPQRPG
						IPPEPPPPGACYKC/HEIRPPGQG
						MPAAQDSS*ATS/LSVRDPTGN
	l				<u></u>	WTVQLTWHPLPEPLELWPKAL

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20379	50747	A	20496	128	302	TAAARHSSRTSSPRSLLQVLEI WPSGQGMPAAWDSS*AASHL WQPLPEP\LELWPKAL
20380	50748	С	20497	80	106	
20381	50749	Α	20498	196	575	
20382	50750	A	20499		505	TKETRFIRGPKTPAPVTDWEGS LPLVFNHCRDASLIIHPGFRGVR PRRDACLSPSPLANLINLTFKVY NNRKKLQFLAFTVRQTSAMSP AHKNFQTPEPQRPGIPPEPPPPG ACYKC/HEIRPPGQGMPAAQDS S*ATS/LSVRDPTGNWTVQLTW HPLPEPLELWPKAL
20383	50751	A	20500	372	563	LRSADLPWEINPLSSCSLLHEKD PPTSSGPQT\TSPRNISPILN/PEK KETRFIRGPKTPAPVMD
20384	50752	A	20501	96	505	WS*QP*GLGSHRHHRRRHHRSS SSSSSSSSSSSKSNSDWQPLPEPL ELWPKAL
20385	50753	A	20502	79	207	
20386	50754	Α	20503	583	736	
20387	50755	A	20504	99	246	DLIFHQRHSVSAESLTMHGLNN *QTSAKRVVQG\TWQPLPEPLE LWPKAL
20388	50756	Α	20505	654	719	
20389	50757	A	20506	71	951	LRSGDLPWEI/NPLSSCSLLREK DPPTTSGPQT\TSPRNISPISNPR/ PKETRFIRGPKTPAPVTDWEGS LPLVFNHSRDTSLIIHPGFRGVR PRRDACLGPSPLAASPTFLGKG PAAPRQTELGPNSSSASAPPPYN PFIASPPHTWSGLQFPSMTSPPP PAQQFTLKKVAGAKGIVKDLIN LTFKVYNNRKKLQFLASTVRQ TPATSPAHKNFQTPELQQPGVP PEPPPRGACYKFQKSGHRAKEC LQPRIPPKPHPICVGPHWKSDCP THLAATPRAPGTLAQGSLTPSQI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20390	50758	Α	20507	1	1010	CSEYEDSSPAPVPATDLSSTLSS
						SVPQPQDTGTSQQLHPLDPWHE
						LLRAQELQGATNHKGYSHAEH
						EHAGLGVQGGNGALAFSNSGH
1						RHAVPTISSGTGRRRTPSSSAFG
						LLNLHQWFVSGFQAFSDRLKA
				1	İ	ALSASLLLRFGDSDWLPSSSAC
						KCLMLGLHFVIVGNICATLKEK
						YSSMLHLDVTMKKNGEKRTRL
ŀ						QKRKKGMPPHPAYEDLNIAAIT
						LPANVVLHQPSGFRTSGQLDPV
						WWSLDTDAHEIWCQDPGLGSG
						DFPWEITPLSSYSLLHEKDPPTT
						SGPQT\TSPRNISPISNPRQRRQV
						LSMDPKLRHRSRTGKAAFPWC
		ŀ				LIIAGTPL
20391	50759	Α	20508	3	539	IVDDCGGAFTMGVIGGGVFQAI
		l				KGFRNAPVCRLLSEAPLFIYSCS
		1				RSVSPTVNVSSERAESRPTLFM
		ľ				AVSLHMAWCLAHIGIRHRLRG
		1				SANAVRIRAPQIGGSFAVWGGL
						FSTIDCGLVRLRGKEDPWNSITS
						GA\LTGAVL\AARSAPLAM\VGS
						AMMGGILL/ALIEGVSILL/TR*T
						ATV
20392	50760	Α	20509	3	778	GEQRAALPPAPKECTHPNVTNT
						LVQAESSDAGVRLRWGRGLTA
						RPDASAMEEYAREPCPWR\IVD
						DCGGAFT\MGVIGGGVFQAIKG
						FRNAPVGIRHRLRGSANAVRIR
						APSTTGSFAVWGGLFSTIDCGL
						VRLRGKEDPWNSIT\SGA\LTGA
						VL\AAP\SGPLAMGGLQQ*WGG
		ļ		+		ILSAPHLRAFGILL\TR*TA\QQF
						RKWAPNSLEDP\SQ\LPPKDVT\P
						APGYP\SYQQYHLRKPLATMGA
				1		TSSVPFPDGSTLKGRAGFPG
20393	50761	A	20510	1	231	LRWRVWSLSLLMFRCVRSFFLL
		1				VGSWSRWFRSEAADLCGECYS
						S*GSTSGVVRFSRWACGLAGLR
						SEAADLAPPTSC
20394	50762	Α	20511	318	494	

SEQ ID	SEQ ID NO:	1	I .		N .	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	1
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
20395	50763	A	20512	<u> </u>	1234	   MNMNIKKIVKQATVLTFTTALL
20373	30703	1	20312	ľ		AGGATQAFAKENNQKAYKETY
						GVSHITRHDMLQIPKQQQNEKY
						QVPQFDQSTIKNIESAKGLDVW
		1				DSWPQQNADGTVAEYNGYHV
		1				VFALAGSPKDADDTSIYMFYQ
						KVGDNSIDSWKNAGRVFKDSD
		ŀ				KFDANDPILKDQTQEWSGSATF
		ļ				TSDGKIRLFYTDYSGSLLRSQM
						PCADLSILKEAAETLDSTSTRNS
						AFLGNTHDEAPIVHVFIPSLAYS
						TNIWSLIFNPTTMIMMCSRDWL
					:	SSCENKRVDLAFGLVYSGTPVQ
		1				KLEQMKFSPQLGCNHFESKDR
						NDFLFAPEEIEKDPQGESLLCDF
						MTKTPKAMATKAKIDKWDLIK
		1				LKSFCTAKETTIRVNKQPTEWE
						1
						KNFAIYPYDKGLISRIYKELKQI
						YEKKIKQPHQKVDERRLLNLTT
						DRLTGEKAYRFIKVQVLRVTQ GMKIQRRARSKELQGDGDEIQR
		İ				WKRLHSPPRLLGEEPHRIIFKLD
						SANDMHKSAPFSSLQSPLWHQP
						VMQEVERQYKEKKDLVREGG
						KCRCQGRPGKVVLTELSLEAR
						GGWMEDIPSRRTSKCGPWELV
						SHCVQNWWVLGLTDFKNEAS
		1				DPRGVKLQTFTVSVTALKAAR
				1		LELLVLPGGLVVSLASGVKLQT
						FAVSVTAHKRSVDPKTEGAGS
						GLGQPRKGLSQCSSGPKG*YPE
20206	50764	<u> </u>	00510	1	521	STKNLNKFTRKKSNNPIKRLMK
20396	50764	В	20513	11120	531	CALODTA AL WEDI SCLAVADA
20397	50765	Α	20514	1139	1562	GALQPTAALWEPLSGLAKARA
						GTLSLQGSVEGVA*AGTGAAC GACGPDGVPGGRGLGGPRTGS
						L .
						SRPALSHFITEEDKDQRVQDQL
						GGVWHRMEIHEQHVGKQQEE
						GDVEDHIPGEDHEGGGEERHIV
20222	50766	<u> </u>	20515		1507	PKQLLVLVDRFLPK
20398	50766	A	20515	1461	1597	
20399	50767	Α	20516	461	1601	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20400	50768	A	20517		914	MNNLRRAALRAVTLTAKVCSF TPEPARPQTHQKEETPNTSEHQ KEQTPDTPPLRTVTLTARVHGFI LEVSETKNPPIPDTPATREAEGL RSRLAVYSTRDSPCVACSGSYT QAQGSLGRKFQDPVSLFEDLTK NTPEPNLRPITSKPVLVQETLSN FALFLKESNGP\VKVWGSH/IKG LT*KALHGF\HVH\EFG\DNTAR LYQVQGPHF\NPLSQKTTVGPK\ DEESHC*DLGQCDCLTKNGVA DVSI*RFCESHSSGDHCIIG\RTL VVPWKKPDDFGQRVGNE\ESSK T\GNAGKSVWPCGVNWIAQ
20401	50769	В	20518	8	173	
20402	50770	С	20519	13	180	
20403	50771	A	20520	3	486	VTETWKCAGIPHRYRSFESC/IW NPVINSDRMDYAGPVCFKNADI GRCAHEKLRED*DGAAPGEIGI QPEADATPYFATPTPVLTHGPC AAESEPALLIGSKQFGLSRNSHI AIAFDDTKVKNRLTIELEVRTE AEFRLLFYMARINHADFATVQL RNGLPY
20404	50772	Α	20521	1	459	
20405	50774	A	20522	2	239	SDSSGSREVDPVAATTMPGAA GVLLLLLSGGLGGVQAQRPQ QQRQSQAHQQRGLFPAVLNLA SNALITTNATCGEKGPEMYCKL VEHVPGQPVRNPQCRICNQNSS NPNQRHPITNAIDGKNTWWQS PSIKNGIEYHYVTITLDLQQVFQ IAYVIVKAANSPRPGNWILERSL DDV\EYKAWQ*H\AVT\DTEAL TP*PN*SPELGHRQNAQD*\EVI CTSFYSKIHPLENGEIHISLINGR PSADDPSPELLE
20406	50774	A	20523	2	445	MODBRANDA
20407	50775	A	20524		2765	MGRRRLLVWLCAVAALLSGA QARGTP\PWRGLR/LPGASRYSL YTTGWRPRLRPGPHKALCAYV VHRNVTCILQEGAESYVKAEY RQCRWGPKCPGTVTYRTVLRP KYKVGYKTVTDLAWRCCPGFT GKRCPEHLTDHGAASPQLEPEP QIPSGQLDPGPRPPSYSRAAPSP HGRKGPGLFGERLERLEGDVQ RLAQTYGTLSGLVASHEDPNR MTGGPRAPAVPVGFGVIPEGLV GPGDRARGPLTPPLDEILS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
ŀ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20408	50776	Α	20525	2	363	RFTKVEMKEK\MLSAAREKGR
	l		1			VTHKGKPISLKADLSAETLQAR
						REWGPIFNILKEKNFQPRISYPA
						KLSFISEQEIKYFTDKQMLRDFV
						ITRPALKELLKEALNMERNNW
						YQPLQKHAKL
20409	50777	Α	20526	1	1566	
20410	50778	Α	20527	1	1604	MAILFEMRLGQQSSAKFPGFQS
						SFFPVIPVNKPSGKFKRDRAHFS
						STRLLWPDCQISQLWGGYLRK
						KGSSPSQGLIDKTPSPWKKAPR
						ERGGCGRSFSRFKHPCLTALKR
						AADLPAQHSSSAKDTSQKGPTD
						TSYRRALAGIWQVPPLGRSFQK
						KDQAAIFAVVQPLQVIPRQTGS
						GVDLQQTSADWQQRGLTVRR
						KTNKQKGIACSLKDPIRGHQHQ
						RPKDHNFSPAREQNWTENEFD
ŀ						KLTEVGFRRWVITNSSELKKHV
						LTQCKDAKNLERSLEELLTRIPS
•						VEKNINDLMELKNTARELREA
					1	YTSFNSPIDQAEERISVIEDQFN
						KINREDKIREKRIKRNKQSLQEI
						WDYVKRPNLQLIGVPESDKEN
						GTKLENTLQNIIQENFPNLAREA
						NIQIQEI\QRTPQRYSLRRATPRH
		ŀ				IISV/RFTKVEMKEK\MLKAVRE
						KGR\VTHKGKPIRLTTADLSAET
						LQARRQSGGQYFNIL*RKGIFQP
	1	ļ				R\ISYPA\KLSFISEGEIKYFIDK\Q
						MLRDF\VTTRDWPALKEHQAC
20411	50779	Α	20528	58	1081	SSMAKPC\GVRLSGEARKQVEV
						FRQNLFQEAE\EFLLR\FLPQKII
						Y\LNPASFQEDFPFNVA*/LLNLS
	1	l				SGAPLGHPPIPRPLHPKDDEM\E
						TDKAGRRKEVP*SVGFFPWGN*
						GKFL\SLVCPGLKPE\VWDFSKR
ĺ						NCILGDLHWIQTPDSPRFEDGN
						DFGGSQSRKKVLEKG*MPVKT
						KV\EAF\QTTISKY/FSSERGDAC
						GPRPPKETHVMDYRALVHERD
						EAAYGELRAMVLDLRAFYAEI
		1				VKSSGIGAHVSFKVQNIHSSYC
						SGPTVSPTCVTGLPASQSPDSG
						AQLASSSGSHTCAAGGAACQS
						RALRLHSSALGWSMGLGTVEQ
						GAALIEEVQAAQEPTEG
	<u> </u>	<u> </u>	L	<u> </u>	L	CHADIDD I YAAYDI IDO

sequence   09/540,217   codon for peptide   of peptide sequence   deletion, \=possise   20412   50780   A   20529   1632   1932   KEGCELCRO   PHHVFPH/V   VNVSQFFIC	=possible nucleotide ible nucleotide insertion)
Sequence	ible nucleotide insertion)
20412 50780 A 20529 1632 1932 KEGCELCRG PHHVFPH/V VNVSQFFIC	
PHHVFPH/V VNVSQFFIC	
PHHVFPH/V VNVSQFFIC	COCRDORDA COLICO
VNVSQFFIC	•
KHVVFGKV	TCQDAKTEWLDC
	KDGMNIVEVMEH
	SNQQEDHHC
	DCSSNP*NYVNRLE
	TAEKF\QALSTGEK
	\FHRIIPGF\TCQSGD
	KSHLPGRNLMTRT
	ILSMANAGPSVNV
SQFFI\CPAK	MPRTEW\LDCKHV
VFGKVKDG	MNIVEVMEHLGSK
NGKISNQQI	EDHHC
20414   50782   A   20531   135   1427   GKPTFASRO	QQPLAEECPASSAL
QRRWLWR	WRWWLRRRRRQR
RSSGSCQGC	QPAGPLGGPPANRG
	MNYNAKDEVDGG
PPCAPGGTA	AKTRRPDNTAFKQQ
RLPAWQPII	TAGTVLPIFFIIGLI
FIPIGIGIFV	<b>FSNNIRE</b> IEIDYTGT
EPSSPCNKC	LSPDVTPCFCTINF
TLEKSFEGN	IVFMYYGLSNFYQ
	RDDSQLNGDSSAL
	YRRNEDKPIAPCG
AIANSMFNI	DTLELFLIGNDSYPI
	AWWTDKNVKFRN
	ERFKGTTKPVNWL
	DPDA*WIHK\DDFI
	LPTFRKLNRLIQRK
	GRYSLNV\NAYYP
	RMIMSAISWMGGK
	FAVGAISFHLGVAL
	NSSNTADITI
20415 50783 A 20532 1 370	10011111111
	FDIWRKKYMRWM
	IDFFRRIDKDQDGK
	LSSKFPTSRLEMSA
	GDGYIDYYEFVAAL
	PITDADKIEDEVTR
	KRFQVEQIGDNKY
	RLVRILRSTVMVRV
1 1 1 1 1	DEFLVKNDPCRAK
	EKFILADGASQGM
	RSRPSSRGASPNRST
	AA\SPQVPATTTPK
	RLPGYLSGKGFHS
	AAARVRTQF\ADS
	SSSSLEAKFGSR\AS
	DFDISEIQSVCSDVE
	PRAGSRPSTAKPS
KIPTPQRKS	PASKLDKSSKR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20417	50785	В	20534	527	4627	
20418	50786	Α	20535	37	315	
20419	50787	A	20536	1314	1577	HGSGSAPTCRWPRTQGRCREIC
						ADGFSGVMRNCTG*EAASGR*
						CKSFVHVYLQLRCACSVCHSA
						GSDSFQLQRSPHTPSTCTHALV
20420	50788	Α	20537	1	1652	MQATKTESGRNGKPGQPSNRC
						HYLLNQGWVANSGKGGRTNIP
						DKELKSGDDPENVHITSFQYLA
						LAAFPRGRARDLQLAMTEPPHP
						SVGSCAARASLTSAAPCSTAPSP
						INHPKAEECGCTARDWQAAPP
						AARVRDPLGEASWAPESEYVC
						NCSVVGSLNVNRCNQTTGQCE
		ļ.				CRPGYQGLHCETCKEGFYLNY
						TSGLCQPCDCSPHGALSIPCNSS
						GKCQCKVGVIGSICDRCQDGY
				1		YGFSKNGCLPCQCNN\RSARCD
				-		ALTGACLNCHENSKGNHCEEC
						KEGFYHSPDATKECLRCPCSAV
						TSTGSCSIKSSELEPECDQCEDG
						YISPNCNKCENGYYNFDSICRK
						CQCHGHVYLISTPKICKPESGEC
						INCLHNTTGFWCENCL*GYVH
						DLEGNCIKKEVILPTPEGSTILVS
						NASLTTSVPTPVINSTF\TPT\TL*
						TIFSVSTSENSTSALADVSWTQF
-						NIIILTVIIIGVVLLMGFEGAGY
						MYRENQNRKLNAPFWTIELKE
						DNISFSSYHDSIPNADVSGLVED
						DGNEVAPNGQMTLTTPIHNYK
20421	50789	Α	20538	1	177	
20422	50790	Α	20539	1	480	
20423	50791	Α	20540	410	896	FSCLKRMHIKAMPEDAKGQDW
						IALVKAAAAAAAKNKTGSKPR
						TSANSNKDKDKDERKWFKVPS
						KKEETSTCIATPDVEKKEDLPTS
						\SETFGLHVENVPKMVFPQPEST
			1			LSNKRKNNQGNSFQAKRARLN
						KITGLLASKAVGVDGAEKKED
						YNETAPMLEQV

SEQ ID			SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence	1	09/540,217	codon for peptide sequence	of peptide sequence	detetion, -possible nucleotide insertion)
1				,		
20424	50792	Α	20541	1	1526	LKDEEDFFDFKAGEEVLARWT
		İ				DCRYYPAKIEAINKEGTFTVQF
						YDGVIRCLKRMHIKAMPEDAK
						GQVKSQHPLSWCCPIDPAGSCN
					ļ	QSMGSEDWIALVKAAAAAAA
		1				KNKTGSKPRTSANSNKDKDKD
						ERKWFKVPSKKEETFNLLLATP
						DVE/RRKEDLPTSS/EKHL*GLH
						VRRNVPKMVFPQPESTLSNKRK
						NNQGNS\FQAKRARLNKITGLL
	1					ASKAVGVDGAEKKEDYNET\A
						PMLEQA\ISPKPQSQKKNEADIS
						SSANTQKPALLSSTLSSGKARS
						KKCKHESGDSSGCIKPPKSPLSP
						ELIQ\VEDLTLVSQLSSSVINKTS
		ŀ				PPQPVNPPRPF\KH\SERRRRSQR
						LGTLP\MPDDSVEKVSSPLSSPL
						D/EGKVFS/LSSSSKISQESSS/IPE
						VPDVA/HICPLEKLGPCLP\LDLS
						RGSEVTAPVAS\DSSYRNECPRA
		1				EKEDTQMLPNPSSKAIADGRGA
		1				PAAAGISKTEKKVKLEDKS\STA
						FGIRSWDFSALLMKIPSYSPISLS
						GLPES
20425	50793	Α	20542	363	1187	FHITMCGICCSVNFSAEHFSQDL
						KEDLLYNLKQRGPNSSKQLLKS
						DVNYQCLFSAHVLHLRGVLTT
						QPVEDERGNVFLWNGEIFSGIK
		1				VEAEENDTQILFNYLSSCKNES
		1				EILSLFSEVQGPW\AFIYYQASS
		1				HYLWFGRDFFGRRSLLWHFSN
		1				LGKSFCLSSVGHPNIWNWQISG
		1				KKFQHLDFFRIDLKSTVISRCIIL
		1				QLYPWKYISRENIIEENVNSLSQ
]						ISADLPAFVSVGSQMEGQTVSL
						EKPVVPLNMDVATSWHWETH
						AVFFPCATYR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	1 ' ' '
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20426	50794	Α	20543	1140	2372	CCHKLHWRLIAVIFPMCHLQER
1						YFKSFLLMYT*KEVIQQFIDVLS
						VAVKKRVLCLPRDENLDSQME
ļ		İ				VLENV**GKANVGNPVFWGGL
						D/CPWLLATLLDRHIPLDEPLVL
				1		LNVAFIAEEKTMPTTFNREGNK
						QKNKCEIPYPKNFSKDVAAAA
						ADSPNKTCQC\PDRITGRAGLK
	-					ELQAVSPSRIWNFVEINVSMEE
	į					LQKLRRTRICHLIRPLDTVLDDS
						IGCAVWFASRGIGWLVAQEGV
		ŀ				KSYQSNAKVVLTGIGCRWSNL
		ŀ				QGYSPSSVSAFKSHGLGRIGIKE
	1					IMMELGRISSLEILGRDDR\VIG
		1				DHGKEARFPFLG*/RMLSPFLNS
						SARFGEKANLDFTPRGIGEKLPF
		1				NALQPVELG\LTASALLPKRAM
		1				Q\FG\$RIAKMEKINEKASDKCG
						RLQIMSLENLSIEKGTKL
20427	50795	В	20544	70	464	
20428	50796	Α	20545	18	552	
20429	50797	Α	20546	1	1257	
20430	50798	В	20547	223	342	
20431	50799	Α	20548	9	5293	KIHRHRLSPATEKGAASSGEGD
						KDPPPPAHEDIAVKGTATAAGT
						GPGTGAAAAAAAAAVPPPHPNI
						RALQTQAPQQIPRGPVQQPLED
		1				RIFTPAVSAVYSTVTQVARQPG
	]	1				TPTPSPYSAHEINKGHPNLAATP
						PGHASSPGLSQTPYPSGQNAGP
}		1				TTLVYPQTPQTMNSQPQTRSPF
		1				FPKGLKIQPP\RATIPNSKSFPFV
		1				PGAQTPTAVYQG*FRHIM\M\V
						NHLPLPYPVPQGPQYCIPQYRH
		1				SGPPYVGPPQ
20432	50800	Α	20549	1	78	
20433	50801	Α	20550	1	2805	MAKGFYISKSLGILGILLGVAA
						VCTIIALSVVYSQEKNKNANSS
		•				PVASTTPSASATTNPASATTLD
						QSKAWNRYRLPNTLKPDSYRV
						TLRPYLTPNDRGLYVFKGSSTV
						RFTCKEATDVIIIHSKKLNYTLS/
						QGHRVVLR\PPDIDKTELVEPTE
						YLVVHLKGSLVKDSQYEMDSE
						FEGELADDLAGFYRSEYMEGN
						VRKVVATTQMQAADARKSFPC
						FDEPAMKAEFNITLIHPKDLTAL
		1				SNMLPKGPSTPLP
20434	50802	A	20551	1	239	
20435	50803	Α	20552	1	654	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20436	50804	A	20553	161	3193	GSLSRLPTITMAKGFYISKSLGI
20.00		-				LGILLGVAAVCTIIALSVVYSQE
		ł				KNKNANSSPVASTTPSASATTN
						PASATTLDQSKAWNRYRLPNT
						LKPDSYQVTLRPYLTPNDRGLY
		i				VFKGSSTVRFTCKEATDVIIIHS
						KKLNFP\LSQGHRVVLRGVGGS
ļ				1		<b>QPPDIDKTELVEPTEYLVVHLK</b>
1						GSLVKDSQYEMDSEFEGELAD
				-		DLAGFYRSEYMEGNVRKVVAT
						TQMQAADARKSFPCFDEPAMK
						ARF\NITLIPPQGT
20437	50805	A	20554	3	429	ind wildinger
20438	50806	Α	20555	3568	4012	VLSTSFPGTELNPPISPLAVNSTE
		ŀ				LLKGELSLSSLGVLSLSDTSVPN
						SVSNSFLLGLRKRLGRGVGISCS
						NALSALDLAACNFSLMRPLME
						NSAKPVLEVEGIGG/TAFDAGA
						GIALSDHFIKLISWYDNEFGYSN
						RVVDLMAHMASKE
20439	50807	A	20556	1	1211	QTAASSFASPGEPHRSNTMGKV
		1	1			KALVNGFDRIGRLVTRAAFNSG
						KGDIVAINDPFIDLNYMVYMFQ
						YDSTHGKFHGTVKAENGKLVI
	1					NGNPITIFQ\ERDPSKIKWG*H\G
						AEYVV\ESTG\VFTTM\EKAGAH
E						FAGGSQKGHSSFAPSTEAIMFM
				İ		MKVNYEKYDNSLKIISNASCTT
						NCLAPLAKVIHDNSGIVERLMI
		İ				TVHVITTTQKTVDGPCRKLRPD
		İ				GHRALQNIIPASTSTAKAMVKV
						<b>IPELNKKLTGMASHVPTAKVLV</b>
						VDPTCH/LGKPAKYYDIKKMM
1						KQASEDPIKGILGYTEHQIVSSD
						FNSDTQSSTFDAGAGITLNDHF
						VK\LISWYDNEFGYSNSVVDLM
						AHVASKELSGYSLKVSFAVQK
						LFSLIRSNLSHAFDAIVFGVFVM
		İ				KSLPVSICRM
20440	50808	A	20557	1	300	PTANVSVVDLTCRL*KPAKYD
						DIKKVVKQASEGPLKGILGYTE
		1				HQVVSSDFNSDTHSSTFDAGAG
		1				IALNDHFVKLISWYDNEFGYSN
1		1		1		RVVDLMAHMASKE
20441	50809	Α	20558	1	3399	
20442	50810	Α	20559	1	3126	
20443	50811	С	20560	298	435	
20444	50812	A	20561	1	888	
20445	50813	В	20562		4536	

SEQ ID			SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first codon for peptide	of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,217	sequence	or peptide sequence	detetion, =possible nucleotide insertion)
	e .					
20446	50814	Α	20563	37	3900	LGLGLSMLVGQGAGPLGPAVV
						TAAVVLLLSGVGPAHGSEDIVV
						GCGGFVKSDVEINYSLIEIKLYT
ŀ						KHGTLKYQTDCAPNNGYFMIP
						LYDKGDFILKIEPPLGWSFEPTT
						VELHVDGVSDICTKGGDINFVF
						TGFSVNGKVLSKGQPLGPAGV
						QVSLRNTGTEAKIQATATQ\PG
						GKFAFFKVLPGDYEILATHPTW
						ALKEASTTVSVTNSNANAASPL
						IVAGYNVSGSVRSDGEPMKGV
						KFLLFSSLVTKED\A
20447	50815	Α	20564	1	310	EEETVPTTAGASPGPPRNKKNR
						ELRPQRPKNAYILKKSRISKKPQ
						VPKKPREWKNPEPSAAAS*GGE
						GQAHGGPAICPGQICALSQTPG
						LPGNSLSPRSPVGK
20448	50816	Α	20565	3	267	
20449	50817	В	20566	267	560	
20450	50818	Α	20567	1	429	LIGLDPEPLAEVDVISLEQGRRK
			!			QIERLGYDPQAKAPFQPKPKQK
						GRSSTASLVKRKRKVMDEE\NR
						VK*TIGNGPGLPKGCTLPACAL
						CPCQPAHFLTLCPVSLPSPVPQD
						KVRQSLQQQHHKEAK/APKPTG
						ARPSALDRFVR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	1 ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' '
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
20451	50819	Α	20568	2	1951	FKPTVWQLRGVLHVDRLFVWA
						SMETAPKPGKDVL\SKKDKLQT
						KRKKP\RRYWEEETVPTTAGAS
						PGPPRNKKNRELRPQRPKNAYI
		İ				LKKSRISKKPQVPKKPREWKNP
		ļ				ESQRGLSGAQDPFPGPAPVPVE
						VVQKFCRIDKSRKLPHSKAKTR
	1					SRLEVAEAEEEETSIQSLLRSEL
		1				LLAEEP\GFLEGEDGEDTAKICQ
		l				ADIVEAVDIASAAKHFDLNLRQ
						FGPYRLNYSRTGRHLAFGGRRG
						HVAALDWVTKKLMCEINVME
		1				AVRDIRFLHSEALLAVAQNRW
		1				LHIYDNQGIELHCIRRCDRVTRL
		1				EFLPFHLLLDTASETGFLTYLDV
		1		:		SVGKIVAALNARAG\RLDVMSQ
					-	NPYNAVIHIGHSNG\TASLWESS
		l				YGREPLAK\ILC\HRGLGSGAVA
						\VDSYRHVTWATSGP*DHQLED
						LFDFARGRY\QPLEHSGPCPMG
						AGHLAFSQRELL/VLAGLGDVV
						NIWAGQGKASPP\SLDRPYFTH
						RV\*GPVHGLQFCPFEDVLGVG
		ŀ				HT*G\ITSML\VPGAGEPNFDGL\
						ESNP\YRSRKQRQ\EWEVKALL
=						EKVPAELICLDPRALAEVDVISL
		İ				ETGKRRKQIERLGYDPQ\AKAP
				:		F\QPKPKQKG\RSSKA\TLVKRK
						RKVMIEDHKG/DKVRQSLQQQ
						HHKEAK\AKPTG\ARPSALDRF
20452	50820	Α	20569	1	437	
20453	50821	Α	20570	157	425	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20454	50822	Α	20571	207	1562	LVIVFFYPQHTAPGPPLTVLNQ
1						AMHARGPHGQLSPALPLASSVL
						MLLMNLAHLGQPWMLGPLTW
						HISPFSLPVHRLLTHALGHTALP
						GLLLSLLLLPTVGWQQECHLGT
r r		1				LRFLHASALLALASGLLAVLLA
		l				GLGLSSAAGSCGYMPVHLAML
ŀ						AGEGHRPRRPRGALPPWLSPW
						LLLALTPLLSSEPPFLQLLCGLL
						AGLACILCLSPGPVGMCGFCG
<b> </b>		ļ				WVPGKEVAQRGEGPSNTKEPC
						THIAWPVAWAHPWLLLVFLDR
						PPCPGQTDAAGAFRWLEPSERR
						LQVLQEGVLCRTLAG\GMPVQ
						VVGTVAGLPVRPSSLP/SSSRPPI
	1	ŀ				PGPPYVASPDLWSHWEDSALPP
		ŧ				PSLRPVQPTWEGSSEAGLDWA
		l				GASFSPGTPMWAALDEQMLQE
						GIQASEVGSPGIGAGVALTLPTS
						PPTPNRLQQLERMGFPTEQAVV
						ALAATGRVEGAVSLLVGGQ
20455	50823	A	20572	460	916	AHSPRDHPWTVLNQAMHARGP
						\HGQ\LSPALPL\ASSVLMLLMST
1						LWLVGAGPGLVLAPELLLDPW
						QVHRLLTHALGHTALPGL\LLS
						LLLLPTVGWQQECHLGTLRFLH
-						ASALLALASGLLAVLLAGLGLS
						SAAGSCGYMPVHLAMLAGEG
1						HRPRRPRGALPPWLSPWLLLAL
						TPLLSSEPPFLQLLCGLLAGLAY
1						AAGAFRWLEPSERRLQV\LQEG
						VLCRTLAGCWPLRLLATPGSLA
					]	ELPVTHPAGVRPPIPGPPYVASP
1						DLWSHWEDSALPPPSLRPVQPT
						WEGSSEAGLDWAGASFSPGTP
		Ì				MWAALDEQMLQEGIQA\SLL\D
						GPAQEPQSAPWLS\KSSVS\*MR
						LQ\QLERMGFP/TTEAAVVALA\
						ATGRVEGA\VSLLVGGQSGT\ET
	1					LVTHGKGGPAHSEGP\GPPWAN
1						CPQHCLLASSVLMLLMSTLWL
						VGAGPGLVLAPELLLDPWQVH
						RLLTHALGHTALPGLAPEPAAP
						AHCGLAAGVPPGHAEIPACLSP
						ARPGFWAAGSAAGRPWAVQCS
						RQLWIHACPPGHAGWGRTPP
20456	50824	Α	20573	174	401	CTVVFSVLAELSCTSRFCPNMW
						TPLREPMTENGRWGLSVRVLE
1						HGGNQSNSACGPEPPEGEMGS*
1		1			1	LEED*LPALIP

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20457	50825	A	20574	450	1029	ATQLPPQPWQTPAALGWAAAT SLMPYRAPSPPARAQHPPALCL APPPPLWLQLPHLEASHSLVEA RPNPPVSTLAQQGIQPSPRHLPR CPSLRPRQQPPQQVP/PQPAAPT PTATITSTGPSLFASIATAPTSSA TTGLSLCTPVTTAGAPTAGTQG FSLKAPGAA\PAPPQQHPPLPPP PPPPPPAAAPPALP
20458	50826	С	20575	115	276	
20459	50827	A	20576	3		ASRLGPREGGARKNPGGPCSAT QYISVPSP\GAASFPVLGSQSLLT TVSPPLPA*P/LAMAL*PRPLCA PGPSPRSPRLRSPLPPSFFSRGRR RLSLLQPWSE
20460	50828	Α	20577	2	388	
20461	50829	С	20578	244	487	
20462	50830	В	20579	5	569	
20463	50831	A	20580	204	676	RTDQYPQRGETEEGLEGQLLPH RQLPGGLGAHRGPDPACGLRD SSRGVPLAGSPLRVR*GRGHDL RQSGGVRPGGGSNRGLVVAGD ARGRAAPLAGGVPAAADRSRL PGVPAAREPHGPTPAPDPDSAA EPGAVSTLRAEPTPQPVVSTGC CSSTWN
20464	50832	В	20581	1	1413	
20465	50833	В	20582	60	224	
20466	50834	В	20583	8	2469	
20467	50835	Α	20584	841	1436	HHQVPLPVGRERGGKRGPGSW ETCLSVLGPAIPPSEPWPVSPSW VHSLVPGEQALPLICSQPPMAP QWPETGPGPHLVHRTCRVCLN WQGMTRTRGYQAGPSSRRTIPT PAQLPAPLASLPPRQDPPSSPAE PPSGRFSMEPWWEPARHPPGSP AAPCSSHRIT/APPTPSLAAQPG/ PVGSNPPLSPEGLCLAGTPACG W
20468	50836	Α	20585	1799	5195	SKGNGSPSTTQTQTPRAAPDGS LSEETPAGPPTCSVPPASALPRQ QYAKSLPVSVPVWGFKEKRTE ARSSDEENGPKPKPASVDANTK LTRSLPCQGSLKRATSCERAGK AVGEEVSGYSLHGCSKERRRP RLKKEGGSGEPGVGYLFISVLV NSNSELIRLINNAIKNDLASRNP TFMCLALHCIANVGSREMGEA FAADIPRILVAGDSMDSVKQSA ALCLLRLYKASPDLVPMGEWT ARVVHLLNDQHMGV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide inscrtion)
				sequence		
20469	50837	İΑ	20586	8	268	SRSSRTWSPIPEKEKTKLQKQR
						EDELIQKIHKLVQKRDFLVDDA
ŀ						EVERLREQEEDKEMADFLRIKL
						KPLDKVTKSPPI*QSQRAHL
20470	50838	С	20587	50	196	
20471	50839	Α	20588	16	1263	SLSSRCMMVAEQGCALEPCQG
						LSHGAALGVLMVVFP/VSSLQQ
						DQLDIIS\MAETT\MMP\EEIELE
						MAKIQRLREVLVRRESELRFMS
						FEEQYNEDHKKGVIDLFTADSL
						KTMTRLVTGVWLAYLAATASN
						PLQEGEHAGEQIGAGGCRSRLE
						GGALTWDPVLLPPVRISRMLLL
						LLIPGELPGIAFKLMNPTKAEPM
						HMFIILEPFDLRKLMDDIQLCK
į						DIMDLKQELQNLVAIPGCTRSM
						MLVSAQLLEGLRKLIIMAEDKG
				,		ADMSRAQSRSKRPPKMLICLLV
						DVFFPEKEKTKL\QKQREG*AN
						PRRFHKLVQKRDFLV\DDAEVE
						RLR\EQE\EDKEMADFLRIK\LKP
						LDKVTKSPASSRAEKKAEPPP\S
						KPTVAKTGLALIKDCCGATQC
						NIHVSPHVGCPGPWGPPPTLLS
20472	50840	Α	20589	31	339	
20473	50841	Α	20590	1	1179	
20474	50842	Α	20591	1	1167	PARSMAATSLMSALAARLLQP
						AHSCSLRLRPFHLAAV/RGNLSL
						GQVGVQCCNHGLLQP*PPGLK*
		ŀ				SSHLSLNEAVVISGRKLAQQIK
			ļ			QEVRQEVEEWVASGNKRPHLS
						VILVGENPASHSYVLNKTRAAA
						VVGINSETIMKPASISE\EELLNL
						INKL\NNDDNVDGLLVQLPLPE
						HIDERRICNAVSPDKDVDGFHV
						INVGRMCLDQYSMLPATPWGV
						WEIIKRTGIPTLGKNVVVAGRS
						KNVGMPIAMLLHTDGAHERPG
]						GDATVTISHRYTPKEQLKKHTI
						LADIVISAAGIPNLITADMIKEG
				1		AAVIDVGINRVHDPVTAG\PGL
						VGNVDFEGVRQKAGYIT\PVPG
		İ				GVGPMTVAMLMKNTIIAAKKG
		L	<u>L</u>	L		AEA*RSEKVLKSKELGVATN

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
20475	50843	A	20592	1	1963	MAASDVNCGACFAELQGNATA
		1				NTPGGTSHHTHLAIHGSCHDSQ
		1				AKEAGRKRQPWPPPDGRSEPAP
		1				DSHPHLSPEALGYFRRALSALK
		l		:		EAPETGEERDLMVHNIMKEVE
		İ				TQALALSTNRTGSEMLQELLGF
						SPLKPLCRVWAALRSNLRTVAC
		l		•		HRCGVHVLQSALLQLPRLLGSA
		1				AEEEEEEEDGKDGPTETLEEL
						VLGLAAEVCDDFLVYCGDTHG
				•		SFVVRTLLQVLGGTILESERARP
						RGSQSSEAQKTPAQECKPADFE
		l		:		VPETFLNRLQDLSSSFLKDIAVF
		l				ITDKISSFCLQVALQVLHRKLA
		İ				QFCAHICNAVIGYLSARGSSVD
		l				GKPLLLFLRDQTSFRLLEQVLL
						VLEPPRLQSLFEEHLQGQLQTL
						AAHPIANFPLQRLLDAVTTPEL
		1				LSPVFEELSPVLEAVLAQGHPG
		1				VVIALVGACRRVGAYQAKVLQ
		l				LLLEAFHCAEPSSRQVACVPLF
						ATLMAYEVYYGLTEEEGAVPA
		i				EHQVAMAAARALGDVTVLGSL
						LLQHLLHFSTPGLVLRSLGALT
						GPQLLSLAQSPAGSHVLDAILTS
		1				PSVT\PSCAAVCLQNLKGTNIVA
		1				LACKSP*QPRVLKCHLEWSSLE
						GP/RRKIAAELGEQNQEL\KEPP
						FGHHVARNVALTTFLKRREAW
		ŀ				EQQQGAVAKRRRALNSILED
20476	50844	Α	20593	55	393	
20477	50845	Α	20594	150	379	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \= possible nucleotide insertion)
	1			sequence		
20478	50846	Α	20595	1	2408	MGVWGIRTPFMVVGLESPVGS
						SPRGQGSCVRLPQRRPDKSHAG
						ARTGPETLRERAQQSNKLPEW
						AEKGSHPARWSPNRGGRWTLP
						TARERRPSRALKARAAGRRQA
						KVADFRLPAAAPPEGAAQLRLE
						SERLPHQDEKEPSSYRGSCGSCP
						VTLRKMPYLGSEDVVKELKKA
						LCNPHIQADRLRYRNVIQRVIR
						YMTQGLDMSGVFMEMVKASA
				+		TVDIVQKKLVYLYMCTYAPLK
	1					PDLALLAINTLCKDCSDPNPMV
						RGLALRSMCSLRMSKLDQWGQ
						AEVLNFLLRYQPRSEEELFDILN
						LLDSFLKSSSPGVVMGATKLFLI
						LAKMFPHVQTDVLVRVKGPLL
		1				AACSSESRELCFVALCHVRQIL
		1				HSLPGHFSSHYKKFFCSYSEPH
						YIKLQKVEVLCELVNDENVQQ
						VLEELRGYCTDVSADFAQAAIF
						AIGGIARTYTDQCVQILTELLGL
						RQEHITTVVVQTFRDLVWLCPQ
						CTEAVCQALPGCEENIQDSEGK
		1				QALIWLLGVHGERIPNAPYVLE
						DFVENVKSETFPAVKMELLTAL LRLFLSRPAECQDMLGRLLYYC
						IEEEKDMAVRDRGLFYYRLLLV
		1				GIDEVKRILCSPKSDPTLGLL\ED
		1				PAERPVNSWASDFNTLVPVYG
		l				KAHWATISKCQGAERCDPKLP
1		1				KTSSFAASGPLIPEENKERVQEL
						PDSGALMLVPNRQLTADYFEK
20479	50847	В	20596	330	2458	
20480	50848	A	20597	2	171	
20481	50849	Α	20598	32	1040	LQLAKLGDWRMQALRHVHCR
			į			FTEG*FMYSL*LTFPPAGYQVT
						GVFMKNWDSL\DEHGV\CTAD
		ŀ				K\DCE\DA\YRVCQIL\DIPF\HQV
						SYGTEVIFFYSSDFLNEYEKGRT
						PNPDIVCNKHIKFSS*SLCSKNL
ļ						TGADAIATGHYARTSLEDEEVF
						EQKHVKKPEGPASWRNCLSVG
			ļ			WFLYTLGQRANIGGLREPWYV
						VEASPPSHSILQAPRTDHPALYR
						DLLRTSRVHWIAEEPPAALVRD
						KMMECPPPRAPLFYPVPCVLTL
						NQDGTVWVTAVTPDARVSPLQ
						FAVFYKGDECLGSGKILRLGPS
						AYTLQKGQRRAGMATESPSDS
						PEDGP\GLSPLL
20482	50850	Α	20599	12	432	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20483	50851	A	20600	2	509	RLHSLLGDYYQAIKVLENIDLS
						TKSMYSRVPEC\QVTTYYYVGF
	1					AYSMMRCYQDAIRVFASILLYI
:						HRTNSMFQRTKYKYEMIIMLN
						EQMHALLAIALTMYPMRI\DESI
						YLHLREKYVDKMLRMQNGDL
						QVYEELFRYSCTKFL*PVPPNY
						DTVHAHYHTESFLQSLRR
20484	50852	В	20601	8	1188	
20485	50853	Α	20602	1	257	
20486	50854	В	20603	210	312	
20487	50855	С	20604	49	288	
20488	50856	Α	20605	13	446	IPVRPTRPADDYESEAAYDPYA
ŀ						YPSDYDMHTGDPKQDLAYERQ
						YEQQTYQVIPEVIKNFIQYFHKT
						VSDLIDQKVYELQASRVSSDVI
			}			DQKVYEIQDIYEN\DAVFLILYK
		1	ŀ			ELYYRHIYAKVSFQSFSQYRCK
						TAKKSEEEIDFL
20489	50857	В	20606	77	611	
20490	50858	A	20607	1	1956	SLVDPRVRPRVRVLASEAAMS
						YPAD\DYESEAAYDPYA\YPSD
						YDMHTGDPKQ\DLAYE\RQYEQ
						\QTYQVIP\EVIKNFIQYFHKTVS
İ						DLIDQK\VDELQASPCLQVIVID
						QERLYEDSRTIY*EQAWTKL\TE
						RFF\KNTPWPGGWKPLVPQGLA
		1				MDA/VSFLDFYPKNYTYRHIYA
		l				KVSGGPSL\EQRFESYYNYCNL
			j			FNYILNADGSRSPLNYPNQW\L
	-	İ				WGYLSMEFIYPVFSHF\SQYRW
						*DLPKKSEEEI\DFLRSNPKIW\N
	1	l				VHSVLNVL\HSLVDKSNINRQL
	1					GGYTQAGGDP*ECGWGSMGR
						HSLYMM\LHYFSLVR/GFSRLHS
	1					LLG\DYYQAIKVLENIELNKKS
	•	1				MYSRVPRVPRSPTYY\YVGFAY
						LMMRRY\QDAIRVFANILLYIQ
						RTKSMF\QRT\TYKYEMI\NKQIE
		1				QMHAL\LAIALT\MYPMR\IDESI
						HLQLREKYG\DK\MLRMQK\GD
		1				PQVYE*LF\SYSCP\RFLAA\VVP
						TYDNVHPN\YHKEP\FLQ\QLKG
						VFLDEVQ\QQGPAFKPIRSFLKL\
						YTTHALLAKPGLASLDPHRGRE
						FRDPRFLCLSNHQG*RNLVLGP
						AGISSPWIGEISVSPQRVDLPTL
	-					NKDMIPHRGTPKVPQAVMGDF
						LHPVRSHKFGGA*SEPLKEGWG
						QRPWMIFTTHFREPVFDVIIGQG
20491	50859	A	20608	2	413	
		<u> </u>		<del></del> _	<u> </u>	<u></u>

SEQ ID	SEQ ID NO:		SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \= possible nucleotide insertion)
20492	50860	A	20609	232	621	
20493	50861	Α	20610	1	3590	MAPRKRGGRGISFIFCCFRNND
						HPEITYRLRNDSNFALQTMEPA
	1					LPMPPVEELDVMFSELVDELDL
						TDKHREAMFALPAEKKWQIYC
	·					SKKKDQEENKGATSWPEFYID
	ŀ	l				QLNSMAAGILEGVFVHIVITHC
		l				LQNQQQLLHDKQGNRHFICCL
						RSHQHPMLPACSNVHLSSFDRK
		l				TLCKDALCPLPASHVSRYLRKS
		l				LLALEKEEEEERSKTIESLKTAL
						RTKPMRFVTRFIDLDGLSCILNF
						LKTMDYETSESRIH
20494	50862	Α	20611	203	3480	SHQEIEQNSAMAPRKRGG\RGI
			1			AFIFCCLRNNDHPEITYRLRNDS
						NFALQTMEPALPMPPVEELDV
						MFSELVDEL\DLPDKHREAMFA
						LPAEKKWQIYCSKKKDQEENK
						GATSWPEFYIDQLNSMAARKSL
		ļ				LALEKEEEEERSKTIESLKTALR
						TKPMRFVTRFIDLDGLSCILNFL
						KTMDYETSESRIHTSLIGCIKAL
		ŀ				MNNSQGRAHVLAHSESINVIAQ
						SLSTENIKTKVAVLEILGAVCLV
						PGGHKKVLQA
20495	50863	Α	20612	1	338	PGGALRVPDAPAGAEGQELCR
						SDFPVGGAGVGDEEGVQCPAP
						AAHRDDTDLRGAH*EVQDAAG
						RRKQPDREQPAGAEEGAPHLPE
		İ				RVPPG*RHGTCTDRGQARACG
						GPLAPE
20496	50864	В	20613	610	6429	
20497	50865	В	20614	28	280	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20498	50866	Α	20615	160	1397	VKMSAQESCLRLIKDFLFVFNL
		ł				FFFALGSLIFC\FGIWILIDKTSFL
į						SFVGL/AAFVPLQIW\SKVL\AIS
						GNLSPWGIALLGLCWGALK\EL
						RCLLG\LYFG\MLLLLFATQITLE
						KSSSSNQR\AQLERSLR\DVVGK
						TI\QKYGINPE\ETAPEEN\WDYV
	1					QSNLRWCGWHYGRDWFKVFIL
						RGNGSEAH\RVTRS*YNFMAT*
i						NSTILNKVIVHQLTRLEHLARSR
						HSAD\ICAVPAESHITRKG\CAQ
						GİQK\WLHNTLIFP*WAICLGVG
		l				LLE\LGFMTLSNIPVEETVDHGL
		ļ				PTGSARYRLGPAPPQSRAPPPVS
						VRWALPVLPVNICLIPRLAVEPS
	:					GVHISPGDTTWLRAPAAVTSPT
						GPGAFVHSFLSPSVGLPPPTRLF
						FTQTSNKPPAFLVNDHKKGGRS:
						KKALEGPNVTRA
20499	50867	Α	20616	290	492	
20500	50868	Α	20617	1164	2134	NNYNННННННННОУУNНН/
						ННҮННННННИҮЙНННН
						ННИУКНИНННҮНОНИНИУИН
						нниннняуунинниннинни
						YNНҮНННННННННКЕҮНН
						НН/ННННҮННННННҮМҮМҮНН
						НҮННННОНҮННИНННИНН
						YNHHY\HHNQYHYHHYHHHD
						YYHHH/HYHHHNYYYLYNRYH
						НҮННҮНQSNНҮКНННННҢQY
						HHN/HEPFA/HHT*ILFCS*KQDR
						NLHCAWPFFFPFFNCHHHCHH
						HHQQQQHSGVKTKQSRPESLLP
						ERCIYLHQKHHPLHRPGAPQHQ
						KTASWRGQAREAAGSSPGSLSS
						HTADHEVEAFKRCFYSRF
20501	50869	В	20618	164	1526	
20502	50870	Α	20619	1	762	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20503	50871	A	20620	69	1054	AISCHVVHAGHQQTQITGNDPF PVVTYSKGPNLQSSRSDMDDST EREQSRLTSCLKKREEMKLKEC VSILPRKESPSVRSYKDGKLLA ATLLLALHACCLTEVSFYQVAA LQGDLASLRAELQGHHAEKLP AGAGAPKAGLGGSSSCSPRDLK IF*TTSSREKAIP/GSEQQK*GVP FQGPEETVTQDCLQLIADSETPT IQKGSYTFVPWLLSFKRGSALE EKENKILVKETGYFFIYGQVLY TDKTY\AMGHLIQRKKVHVFG DELS\LVTLFR\CIQNMPETLPNN SCYSAGIAKLEEGDELQLAIPRE NAQISLDGDVTFFGALKLL
20504	50872	A	20621	3	855	LQNSHGHTMIRTLLLSTLVAGA LSCGDPTYPPYVTRVVGGEEAR PNSWPWQVSLQYSSNGKWYHT CGGSLIANSWVLTA\AHCISSS\R TYRVGLGRHNLYVA\ESGSLA/ MSSVSKIVVHKD\WNFNQISKG\ NDI\ALLKLANPVSLTDKIQ\LA\ CFPPAGTILPN\NY\PC*VTGLGE SCRTNG\AVPDVL\QQGR\LLAV DYATCS\SSA\WWGSSVKTSMIC A\GGDGVISSCNGDSGGPLNCQ A\SDALWQVHGIVSFGSRLGCN YYHKPSVFTRVSNYIDWINSVI AYN
20505	50873	A	20622	19	414	SVCWEDRYLKARMEESPLSRA PSRGGVNFLNVARTYIPNTKVE CHYTLPPGTMPSASDWIGIFKP/ GYLPKPGAQLYQFRYVNRQGQ VCGQSPPFQFREPRPMDELVTL EEADGGSDILLVVPKATVLQNQ
20506	50874	A	20623	1	393	AKSWQEEQSAQAQRLKDKVA QMKDTLGQAQQRVVSEAPWQ QETGNSKGKL*RSLRV*REGER HG*RLNNTTPSPNAVEWRER/IT SLADE*FTRNCDLHLLLSPEAEL EPLKEQLRGAQELAASSQQKAT LY
20507	50875	A	20624	56	416	NRPLLALVASDTQPLHTHTHTH TPLSDAPKPIPGA/HLP/CSYLEF PLTSVPHPSLFHNLRYSVWRGF LYF*GFLYSVSSPRISSHGKK*CI CAFCEEWGEQVVPGIPISKAPLP LQVPPQQ
20508	50876	Α	20625	115	392	
20509	50877	В	20626	1	3388	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20510	50878	A	20627	3	1889	SSPREKTSDSSHRPSRHGFLFLR LVGLSPFSYLCVPPSRPVPGSPR SLSAMRLLPLAPGRLRRGSPRH LPSCSPALLLLVLGGCLGVFGV AAGTRRPNVVLLLTDDQDEVL GGMTPLKKTKALIGEMGMTFS SAYVPSALCCPSRASILTGKYPH NHHVVNNTLEGNCSSKSW\QKI QEPNTFPAILRSMCGYQTFFCR GNI*MKYGAP\DAGG\LG\HVPL GLGVTWYALEKNS\KYYNYTL FYSMGKARKHGENY\SVDYLT
	60070		20/09		0.52	DVLANVSLDF\LDYKSNF*PFF MMIATP\APHSPWTAAPQYQKA FQNVFAPRNKNFNIHGTNKHW LIRQAKTPMTNSSIQFLDNAFR KRWQTLLSVDDLVEKLVKRLE FTGELNNTYIFYTSDNGYHTGQ FSLPIDKRQLYEFDIKVPLLVRG PGIKPNQTSKMLVANIDLGPTIL DIAGYDLNKTQMDGMSLLPILR GASNLTWRSDVLVEYQGEGRN VTDPTCPSLSPGVSQCFPDCVCE DAYNNTYACVRTMSALWNLQ YCEFDDQEVFVEVYNLTADPD QITNIAKTFDP\ELLGKMNYRL MMLQSCSGPTCRTPGVFDPGY RFDPRLMF\SNRGSVRTRRFSQT SSCSDPHTASADGSLHALFLMK
20511	50879	Α	20628	132	253	
20512	50880	A	20629	1	1269	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20513	50881	A	20630	1	1920	MDEAEFSSFVVRTNTCGELRSS
		l				HLGQEVTLCGWIQYRRQNTFL
						VLRDFDGLVQVIIPQDEVIENFL
						LLSKSFFDAIAGSAASVKKILCE
		}		}		APVESVVQVSGTVISRPAGQEN
						PKMPTGEIEIKVKTAELLNACK
		-				KLPFEIKNFVKKTEALRLQYRY
						LDLRSFQMQYNLRLRSQMVMK
		ŀ				MREYLCNLHGFVDIETPTLFKR
						TPGGAKEFLVPSREPGKFYSLP
		İ				QSPQQFKQLLMVGGLDRYFQV
						ARCYRDEGSRPDRQPEFTQIDIE
						MSFVDQTGIQSLIEGLLQYSWP
						NDKDPVVVPFPTMTFAEV\RPP
						YGTDKPDTRFGMKIIDISDVFR
				}		NTEIGFLQDALSKPHGTVKAICI
			1			PEGAKYLKRKDIGSIRNFAADH
		1				FNQEILPVFLNANRNWNSPVAN
						FIMESQRLELIRLMETQEEDVV
		ĺ				LLTAGEHNKACSLLGK\LRTGN
						VAGPSRKQEGVVLRDPTLFSFL
:			İ			WVVDFPLFPGQRRKIQESWDR
						GQHPF\TAPHPR*IT\LLYTEPKK
						ARSQHYDLVLNGNEIGGGSIRI
						HNAELQRYILATLLKEDVKMLS
ļ				1		HLLQALDYGA\PPHGGIALGLD
						RLICLVTGSPSIRDVIAFPKSFRG
				1		HDLMSNTPDSVPS*GTEALSYP
		1				SLQANRLQSRKSSLNHAYHAES
20514	50882	Α	20631	1	425	KEPVGCVNNISFLASLAGSTSR
						NRLQSTRGAGRLQNSGTGLSTN
		1				LQHFQEENFRKSSPQLEHT/WSF
						FGECTTHHLPPVKAPLQTKKKT
						TNHCFLCGKKTGLASSYECRQ
						VGELEKLRCFLCDNPAWGLCKI
						KYLNCLFITSL
20515	50883	Α	20632	1	1647	
20516	50884	A	20633	1	3096	
20517	50885	Α	20634	1	419	
20518	50886	Α	20635	558	1070	SFRFGHSHGLWKRKKVA*GSN
		1				AIQHNMLQGKQNFHGAALWY
						KGNSTQSKRPQRAP*ESSTTKR
						GLQLSRQTLQLDPGGTVPCSEE
		1				YCICQ*EGPASRNRYNGVPSHF
				1		PSPTCIRESSHSGKILKSRLRCW
						GWPTDPWSVLPMDPRTSPLCA
		<u>L</u> _	L	<u></u>		AGSQAVVAHSGRQQKYPAL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide sequence		in USSN 09/540,217	location of first codon for peptide		*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
20519	50887	Α	20636	3887	4573	KKTTLLTLSKTRNILNPFKVVK KKAYGSRKHLRQ/HLAINPAAL
						LPTAASQISEVKPVLPELAFPSS
						EHRRSHGLESVPVLPGSGEAGV SFDLPAQADTLHSANKSRVKM
						RGKRRPQTRAARRLAAQESSEA
						EDM\RSPEDPLHSGLMAPFPQM
						AIGHSSEQPVEKTALRRPWQLP
						LHLGKVVLCLEWTQAPLQSLW
						VIPEGRLTFLILGTFFPRALDLSP
						WREQNPRQR
20520	50888	Α	20637	1	2988	MATAPSPPMLTLAPAWLCPAT
		l				TTPRDHSEASLEALPGDPPAHP
						AAAATAAAVIPALEPPHTPVYA
						IFVNHSLLDLYIDRPLPYLIGSK
		ļ				LFMEQEDVGLGELSSEEGNADE
		Ì				QSSGLEGACKVNLELLLTNAD
						KGLKAPEQHLKHVAGVLNGES
						VETSVLNYRELSPHKNRLLSPL
						RCSAPMSLHNSLVKPERQSKCF
						EFGKLQPSSSQSLDVQNITDSSF SRTTCFQGVKVDSLGKRSDVIS
i		ŀ				KVEARDITEMTNKA
20521	50889	A	20638	1	1176	KVEARDITEMITY
20522	50890	A	20639	2	367	
20523	50891	Α	20640	124	441	RYPRVPENRCFSELLFFQAPLG
						VGMARGNQR\ELARQKNLKET\
						QEI\SRGKGKEDSLTALSRKQRD
}				}		S\*IIAKKQKAA\NEKKS\MQTRE
			ļ			NVMTGYLENPGATAQLGVS
20524	50892	A	20641	32	232	
20525	50893	A	20642	1	256	
20526	50894	C	20643	133	509	OKDODI EKIH KANA MI MAKADA
20527	50895	Α	20644	88	685	QKRGRLFKIILKVVLYLMKVIVI HRKMFQRAQELRWRAEDYHK
						CKAFSDHPLCKKASLQLGQNG
		1				GSGASSFFRIALLALPHS*NFKK
						QDGPPATSSNSTTFYN**LPEPQ
						DTSRMSSYSLSTLSSSTLRG**S
						QDSSLSYSGGRGRKITETQEAE
				-		VE*AENRSLHSSLGNKKKSVSG
						TKEKNQKTKQNPHSKNKLTKN
20528	50896	Α	20645	42	946	
20529	50897	С	20646	1	453	
20530	50898	С	20647	771	1223	
20531	50899	С	20648	324	659	
20532	50900	С	20649	427	654	<u> </u>
20533	50901	В	20650	343	1774	
20534	50902	В	20651	23	525	
20535	50903	В	20652	2354	5689	
20536	50904	В	20653	448	3506	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		-		sequence		
20537	50905	В	20654	12	182	
20538	50906	C	20655	210	422	
20539	50907	c	20656	145	357	
20540	50908	В	20657	1	327	
20541	50909	В	20658	1	2595	
20542	50910	В	20659	122	472	
20543	50911	В	20660	163	418	
20544	50912	C	20661	86	517	
20545	50913	В	20662	334	2946	
20546	50914	В	20663	11	700	
20547	50915	В	20664	184	2372	
20548	50916	C	20665	267	467	
20549	50917	В	20666	131	268	
20550	50918	В	20667	18	255	
20551	50919	В	20668	100	356	
20552	50920	В	20669	120	839	
20553	50921	В	20670	355	504	
20554	50922	В	20671	1	1524	
20555	50923	В	20672	175	366	
20556	50924	В	20673	436	792	
20557	50925	В	20674	94	1347	
20558	50926	В	20675	865	1189	***
20559	50927	В	20676	241	423	
20560	50928	В	20677	124	352	
20561	50929	В	20678	1	847	
20562	50930	В	20679	262	4624	
20563	50931	C	20680	76	297	
20564	50932	В	20681	1	702	
20565	50933	В	20682	1	1347	
20566	50934	В	20683	78	2874	
20567	50935	B	20684	133	7317	
20568	50936	В	20685	294	1449	
20569	50937	C	20686	139	367	
20570	50938	c	20688	154	445	***
20571	50939	В	20689	1	1086	
20572	50940	В	20690	111	440	
20573	50941	В	20691	303	462	
20574	50942	В	20692	1	2905	
20575	50943	С	20693	349	453	
20576	50944	C	20694	233	376	
20577	50945	В	20695	1	2016	
20578	50946	В	20696	1464	3022	
20579	50947	В	20697	1	1059	
20580	50948	В	20698	i	933	
20581	50949	В	20699	1	930	
20582	50950	C	20700	129	215	
20583	50951	C	20701	283	408	
20584	50952	В	20702	1	1134	
20585	50953	В	20703	1	525	
20586	50954	В	20704	146	556	
20587	50955	В	20705	1	675	
20587	30955	$I_R$	20705	1	6/5	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
20588	50956	В	20706	1	1107	
20589	50957	В	20707	1	774	<del></del>
20590	50958	C	20708	52	156	
20591	50959	C	20709	1	420	
20592	50960	В	20710	1	1347	
20593	50961	C	20711	1	411	
20594	50962	В	20712	169	248	
20595	50963	В	20713	63	1321	
20596	50964	В	20714	1	359	
20597	50965	C	20715	1195	1674	
20598	50966	В	20716	964	1290	
20599	50967	В	20717	51	389	
20600	50968	В	20718	108	232	
20601	50969	В	20719	425	3545	
20602	50970	C	20720	35	226	
20603	50971	В	20721	1	387	
20604	50972	В	20722	1	1926	
20605	50973	В	20723	523	1242	
20606	50974	В	20724	42	2346	
20607	50975	В	20725	63	287	
20608	50976	В	20726	59	200	
20609	50977	В	20727	1	1176	
20610	50978	В	20728	1	769	
20611	50979	В	20729	1	2009	
20612	50980	В	20730	219	3045	
20613	50981	В	20731	195	1739	
20614	50982	C	20732	126	435	
20615	50983	В	20733	2752	4108	
20616	50984	В	20734	1	1104	
20617	50985	В	20735	24	230	
20618	50986	В	20736	1	1292	
20619	50987	В	20737	1	7977	
20620	50988	В	20738	157	2627	
20621	50989	C	20739	165	533	
20622	50990	В	20740	35	87	
20623	50991	C	20741	68	202	
20624	50992	В	20742	83	2670	
20625	50993	В	20743	162	276	
20626	50994	В	20744	104	245	
20627	50995	В	20745	66	2768	
20628	50996	В	20746	14	3115	
20629	50997	C	20747	69	203	
20630	50998	В	20748	1	2152	
20631	50999	C	20749	114	387	
20632	51000	C	20750	56	250	
20633	51001	В	20751	53	3398	
20634	51002	В	20752	642	2386	
20635	51002	В	20753	1	1242	
20636	51003	В	20754	51	601	
20637	51005	В	20755	192	463	-
20638	51006	В	20756	125	1254	
20038	00015	ո	20730	123	1234	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence	i	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20639	51007	С	20757	208	333	
20640	51008	В	20758	155	4443	
20641	51009	В	20759	1	2529	
20642	51010	В	20760	63	321	
20643	51011	В	20761	226	645	
20644	51012	В	20762	1	1395	
20645	51013	В	20763	258	475	
20646	51014	C	20764	126	263	
20647	51015	В	20765	1	1429	
20648	51016	В	20766	51	3256	
20649	51017	c	20767	83	403	
20650	51018	В	20768	1	621	
20651	51019	B	20769	1	463	
20652	51020	В	20770	1002	3594	
20653	51020	C	20771	107	403	
20654	51021	В	20772	51	401	
20655	51022	С	20773	46	398	
20656	51023	В	20774	1	513	
20657	51025	В	20775	<u> </u>	531	
20658	51026	C	20776	1	423	
20659	51027	В	20777	52	670	
20660	51027	С	20778	43	375	
20661	51028	В	20779	1	1629	
20662	51029	В	20780	1	1713	
20663	51031	В	20781	1	5154	
20664	51031	В	20782	37	585	
20665	51032	В	20783	1	3789	
20666	51034	С	20784	149	367	
20667	51035	c	20785	89	234	
20668	51036	В	20786	113	278	
20669	51037	В	20787	1	1632	
20670	51037	В	20788	1	762	
20671	51039	В	20789	230	3411	
20672	51040	В	20790	95	1150	
20673	51040	В	20791	1	1230	
20674	51041	В	20791	74	3123	
20674	51042	В	20792	1	1719	
20675	51043	С	20794	206	402	
20676	51044		20794	302	424	
	51045	B B	20796	60	80	
20678						
20679	51047	В	20797	1	1248	
20680	51048	В	20798	8	1910	
20681	51049	В	20799	124	807	
20682	51050	В	20800	43	618	
20683	51051	В	20801	7	328	
20684	51052	В	20802	244	385	
20685	51053	В	20803	850	970	
20686	51054	С	20804	97	360	
20687	51055	В	20805	551	3159	
20688	51056	В	20806	861	1110	
20689	51057	В	20807	1	876	<u></u>

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20690	51058	В	20808	T1	849	
20691	51059	В	20809	1	1443	
20692	51060	В	20810	508	2398	
20693	51061	В	20811	53	185	
20694	51062	В	20812	50	633	
20695	51063	В	20813	1	444	
20696	51064	В	20814	58	262	
20697	51065	В	20815	27	1005	
20698	51066	В	20816	1	4860	
20699	51067	В	20817	1	3489	
20700	51068	C	20818	52	363	
20701	51069	C	20819	347	499	
20702	51070	В	20820	1	4140	
20703	51071	В	20821	85	3930	
	51072	В	20822	848	3116	
20705	51073	В	20823	41	1211	
20706	51074	В	20824	248	4307	
20707	51075	В	20825	67	3180	
20708	51076	С	20826	229	467	
20709	51077	В	20827	71	1827	
20710	51078	В	20828	1	1045	
20711	51079	В	20829	1279	1479	
20712	51080	В	20830	1	2460	
20713	51081	В	20831	62	3691	
20714	51082	В	20832	1	1062	
20715	51083	В	20833	1	873	
20716	51084	В	20834	1	1041	
20717	51085	В	20835	161	5445	
20718	51086	В	20836	199	343	
20719	51087	С	20837	86	229	
20720	51088	c	20838	1	225	
20721	51089	Ċ	20839	34	165	
	51090	c	20840	238	455	
	51091	C	20841	200	322	
20724	51092	В	20842	59	164	
20725	51093	c	20843	1353	1793	
20726	51094	В	20844	1	1329	
20727	51095	C	20845	43	324	10-10-11-11-11-11-11-11-11-11-11-11-11-1
20728	51096	C	20846	223	435	
20729	51097	C	20847	212	307	
20730	51098	C	20848	386	529	
20731	51099	В	20849	300	995	
20732	51100	В	20850	876	3312	
20733	51101	В	20851	153	306	
20734	51102	В	20852	39	3131	
20735	51103	В	20853	29	136	
20736	51104	В	20854	112	333	
20737	51105	В	20855	31	474	
20738	51106	В	20856	26	1716	
				27	1234	
20739	51107	B	20857	147	11234	

SEQ ID	SEQ ID NO:					Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/340,217	sequence	peptide sequence	government of the second of th
20741	51109	В	20859	11	1014	
20741	51110	В	20860	128	837	
20742	51111	В	20861	128	931	
20743	51112	В	20862	357	4629	
20745	51113	В	20863	1	1017	
20745	51114	В	20864	1	1908	
20746	51115	В	20865	224	9448	
20747	51116	В	20866	1	1209	
20748		_		138	359	
	51117	В	20867		1567	
20750	51118	В	20868	226	9033	<u>'</u>
20751 20752	51119	В	<u> </u>		2689	
	51120	В	20870	20		
20753	51121	В	20871	1602	2647	
20754	51122	C	20872	72	452	
20755	51123	C	20873	64	423	<u> </u>
20756	51124	В	20874	33	363	
20757	51125	В	20875	131	1675	
20758	51126	В	20876	37	219	
20759	51127	В	20877	1	564	
20760	51128	В	20879	1	651	
20761	51129	В	20880	44	424	
20762	51130	В	20881	28	202	
20763	51131	В	20882	1	3403	
20764	51132	В	20883	110	970	
20765	51133	В	20884	70	485	
20766	51134	В	20885	482	2943	
20767	51135	В	20886	469	1480	
20768	51136	В	20887	156	308	
20769	51137	C	20888	218	348	
20770	51138	В	20889	1	5130	
20771	51139	В	20890	21	1139	
20772	51140	В	20891	1	1302	
20773	51141	В	20892	1	1028	
20774	51142	В	20893	1	858	
20775	51143	В	20894	1	2001	
20776	51144	В	20895	1	546	
20777	51145	В	20896	1	1167	
20778	51146	В	20897	1	1299	
20779	51147	В	20898	1	1260	
20780	51148	В	20899	1	771	
20781	51149	C	20900	155	337	
20782	51150	В	20901	1	1251	
20783	51151	В	20902	106	543	
20784	51152	В	20903	57	2279	
20785	51153	В	20904	905	1120	
20786	51154	В	20905	145	2277	
20787	51155	В	20906	1	1384	
20788	51156	В	20907	1	4775	
20789	51157	С	20908	99	251	
20790	51158	В	20909	1	693	
20791	51159	В	20910	77	2173	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20792	51160	В	20911	140	313	
20793	51161	В	20912	1	392	
20794	51162	С	20913	1	222	
20795	51163	В	20914	1	951	
20796	51164	В	20915	45	1244	
20797	51165	В	20916	166	2544	
20798	51166	С	20917	279	449	
20799	51167	В	20918	1	1250	
20800	51168	В	20919	229	2094	
20801	51169	В	20920	94	434	
20802	51170	С	20921	253	336	
20803	51171	В	20922	1	1314	
20804	51172	В	20923	6	327	
	51173	В	20924	163	3060	
20806	51174	В	20925	20	385	
20807	51175	В	20926	22	58	
20808	51176	С	20927	141	287	
20809	51177	В	20928	37	1063	
20810	51178	С	20929	261	401	
20811	51179	В	20930	66	3148	
20812	51180	В	20931	223	4149	
20813	51181	C	20932	21	278	
20814	51182	В	20933	158	1283	
20815	51183	В	20934	8	896	
20816	51184	В	20935	240	387	
20817	51185	C	20936	312	512	
20818	51186	В	20937	451	611	
20819	51187	С	20938	160	428	
20820	51188	В	20939	110	294	
20821	51189	В	20940	121	1764	
20822	51190	В	20941	1	423	
20823	51191	С	20942	74	316	
	51192	C	20943	136	255	
20825	51193	С	20944	79	381	
20826	51194	В	20945	282	1180	
20827	51195	В	20946	63	371	
20828	51196	C	20947	193	672	
20829	51197	В	20948	85	898	
20830	51198	В	20949	35	1055	
20831	51199	В	20950	1	870	
20832	51200	В	20951	103	1189	
20833	51201	В	20952	501	594	
20834	51202	В	20953	39	1694	
20835	51203	В	20954	489	4797	
20836	51204	C	20955	31	153	
20837	51205	В	20956	259	699	
20838	51205	В	20957	4	231	
20839	51207	В	20958	50	520	
20840	51207	С	20959	322	456	
. / 11/4/11	1-1-00	ı~	20000			
20840	51209	В	20960	163	1152	

SEQ ID NO: of peptide sequence			Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
51211	В	20962	210	3519	
51212	С	20963	34	294	
51213	С	20964	301	381	
51214	В	20965	71	675	
51215	В	20966	1	873	
51216	В	20967	118	1618	
51217	В	20968	424	2415	
51218	C	20969	106	231	
51219	В	20970	71	756	
51220	В	20971	1	1049	
51221	С	20972	94	327	
51222	В	20973	339	2589	
51223	В	20974	89	273	
51224	В	20975	1	2514	
51225	В	20976	472	5831	
51226	C	20977	247	300	
51227	В	20978	1	2487	
51228	В	20979	55	243	
51229	С	20980	207	377	
51230	В	20981	642	4117	
51231	В	20982	44	2127	
51232	В	20983	48	1127	
51233	В	20984	1	3411	
51234	В	20985	74	469	
51235	В	20986	39	1251	
51236	В	20987	223	378	
	51211 51212 51213 51214 51215 51216 51217 51218 51219 51220 51221 51222 51223 51224 51225 51226 51227 51228 51229 51230 51231 51232 51233 51234 51235	of peptide sequence         hod sequence           51211         B           51212         C           51213         C           51214         B           51215         B           51216         B           51217         B           51218         C           51219         B           51220         B           51221         C           51222         B           51223         B           51224         B           51225         B           51226         C           51227         B           51228         B           51230         B           51231         B           51232         B           51233         B           51234         B           51235         B	of peptide sequence         hod in USSN 09/540,217           51211         B 20962           51212         C 20963           51213         C 20964           51214         B 20965           51215         B 20966           51216         B 20967           51217         B 20968           51218         C 20969           51219         B 20970           51220         B 20971           51221         C 20972           51222         B 20973           51223         B 20974           51224         B 20975           51225         B 20976           51226         C 20977           51227         B 20978           51228         B 20979           51229         C 20980           51230         B 20981           51231         B 20982           51232         B 20983           51234         B 20985           51235         B 20986	sequence         09/540,217         codon for peptide sequence           51211         B         20962         210           51212         C         20963         34           51213         C         20964         301           51214         B         20965         71           51215         B         20966         1           51216         B         20967         118           51217         B         20968         424           51218         C         20969         106           51219         B         20970         71           51220         B         20971         1           51221         C         20972         94           51221         C         20972         94           51222         B         20973         339           51223         B         20974         89           51224         B         20975         1           51225         B         20976         472           51226         C         20977         247           51229         C         20980         207           51230         B <td>of peptide sequence         hod 99/540,217         in USSN codon for peptide sequence         codon for peptide sequence         codon for last amino acid of peptide sequence           51211         B         20962         210         3519           51212         C         20963         34         294           51213         C         20964         301         381           51214         B         20965         71         675           51215         B         20966         1         873           51216         B         20967         118         1618           51217         B         20968         424         2415           51218         C         20969         106         231           51219         B         20970         71         756           51220         B         20971         1         1049           51221         C         20972         94         327           51222         B         20973         339         2589           51223         B         20974         89         273           51224         B         20975         1         2514           51227</td>	of peptide sequence         hod 99/540,217         in USSN codon for peptide sequence         codon for peptide sequence         codon for last amino acid of peptide sequence           51211         B         20962         210         3519           51212         C         20963         34         294           51213         C         20964         301         381           51214         B         20965         71         675           51215         B         20966         1         873           51216         B         20967         118         1618           51217         B         20968         424         2415           51218         C         20969         106         231           51219         B         20970         71         756           51220         B         20971         1         1049           51221         C         20972         94         327           51222         B         20973         339         2589           51223         B         20974         89         273           51224         B         20975         1         2514           51227

NO: of peptide hod in USSN location of first codon for last amino ac sequence of peptide sequence	1
	1111
	deletion, \=possible nucleotide insertion)
sequence	
20869 51237 A 20988 1 2577	MPPNRHRMRGETTLWNPGCDH
	AGIATQVVVEKKLWREQGLSR
	HQLGREAFLQEVWKWKEEKG
	DRIYHQLK\KLGSSLDWDRACF
	TMDPKLS\AAVTEAFVGLHEKG
	IIYRSTAFVNWPCTLNSAISDIE
	VDKKELTGRTLLSVPGYKEKV
	EFGVLVSF\AYK\VQGSD*/DDE
	EVVVANKLGSKT\MLGDVAVA
	VPPPKDTR\YQHL\KGKNVIHP\F
	LSRSLSPLSSDEFVDMDF\GTGA
	VKDHPPRHDQNDYEVGQRHGL
	EAISIMDSRGA\PIKCASAFPGPA
	AGLRPGKRCWWR*RNRGLFRG
	IEDNPMVVPLCNRSKDVVEPLL
	RPQWYVRCGEMAQAASAAVT
	RGDLRILPEAHQRTWHAWMD
	NIREWCISRQLWWGHRIPAYFV
	TVSDPAVPPGESEDLSVFYPGT
	LLETGHDILFFWVARMVMLGL
	KLTGRLPFREVYLPCPSCENAH
	GRK\MSKSLGNVIDPLDVHL/Y
	GISLQGLHNQLLNSNLDPSEVE
	KAKEGQKADFPAGVPECGTDA
	LRFGLCAYMSQGS*K*TWNVN
	RIMG*RQFCNKLWKCTKFALR
	GLGKGFVPS\DTSNSAGHESLA
	DR*IRNLLTEAVRLSNQGFQAY
	DFPAVTTAQYSFWLYELCDVY
	LECLKPVLNGVDQV/AQAECAR
	QTLYTLPWTLGLRLLSPFMPFV
	TEELFQRLPRRMPQAPPSLCVT
20870 51238 A 20989 2 4083	SQRGEHVGAGCRGGDLRELGS
	PGEPPTVSARSTPPPGPLGGSQP
	RPAPGCEVDSPAQSDHLPPVFP
	LRSDLLITMSTLYVSPHPDAFPS
	LRALIAARYGEAGEGPGWGGA
	HPRICLQPPPTSRTSFPPPRLPAL
	EQGPGGLWVWGATAVAQLLW
	PAGLGGPGGSRAAVLVQQWVS
	YADTELIPAACGATLPALGLRS
	SAQDPQAVLGALGRALSPLEE
	WLRLHTYLAGEAPTLADLAAV
	TALLLPFRYVLDPPAR

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20871	51239	A	20990		756	MDMGAIKEGESRRPGDRMTVA GTKLLHLQSTHLDFISLSGAPSG ALELSKAQPAHAQDTRVLMKM VIIWDLSQVPLQPGHDGHSFRIH DYLYFQVLSPGDIRYIFTATPAK DFGGIFHTRYEQIHLVPAEPPEA CGELSNGFFIQDQIALVERGGCS FLSKTRVVQEHGGRGGDHL*Q RS*QMTASTW/QMIQDSTQRTA DIPALFLLGRDGYMIRRSLEQH GLPWAIISIPVNVHQHSPPLSCC NRPFLFW
20872	51240	Α	20991	2	393	
20873	51241	A	20992	1066	1824	
20874	51242	Α	20993	3	902	
20875	51243	Α	20994	1	2163	
20876	51244	A	20995	1	1890	
20877	51245	A	20996		2882	MKQQLPVSYLQVIFNHWASRIS GARGAGWGPVRLGRLCADRAL DGESRPGGGGQSGVPASEASQ KAAHGRRLPVPARLLRCAHSA LEPGRGSASSPRNSTTSDNDQPP DYSFSKRCDSHSLGFSLVWESR NLTLPPTSKLLHFSAFHFFIYKC RAEGHKLYKDLKNFLSAVKVM HESSKRVSETLQFIYY\SEWYGH EELKAIVWNNDLLWEDYEEKL ADQAVRTMEIYVAQFSEIKERI AKRGRKLVDYDSARH

2650

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20878	51246	Α	20997	1	1791	MPGARTSSSGASENHRARGQG
						GGPQGVGRMAEGKAGGAAGL
						FAKQVQKKFSRAQEKVLQKLG
						KAVETKDERFEQSANNFYQQQ
						AEGHKLYKDLKNFLSAVKVMH
						ESSKRVSETLQEIYSSEWDGHE
						ELKAIVWNNDLLWEDYEEKL
		ŀ				ADQAVRTMEIYVAQFSEIKE\RI
						AKRGRKLVDYDSARHHLEAVQ
	İ					NAKKKDEAKTAKAEEEFNKAQ
		ł				TVFEDLNQELLEELPILYNSRIG
						CYVTIFONISNLRDVFYREMSK
						LNHNLYEVMSKLEKQHSNKVF
						VVKGLSSSSRRSLVISPPVRTAT
			ŀ			VSSPLTSPTSPSTLSLKSESESVS
						ATEDLAPDAAQGEDNSEIKELL
					•	EEEEIEKEGSEASSSEEDEPLPA
				•		CNGPAQAQPSPTTERAK\SQEE
		1				VLPSSTTPSPGGALSPSGQLSSS
		1				ATEVVLRTRTASEGSE\RPKKT
		İ				ASIQRTSAPPRRPPPPRATASPRP
		1	[			CSGNIPSSPTASGGGSA\TSPRAS
						LGTGTASPRTSLEI\*PNPE\PPEK
						PVRTPEAKENENIHNONPEELC
		1				TSPTLMTSQVASEPGEAKKME
		1				DKEKDNKLISADSSEGQDQLQV
		1				SMVPENNNLTAPEPQEEVSTSE
20879	51247	Α	20998	3	273	RRPRQRPTDQTQWFSILPDFSLD
						LQEGPSVESQT\HSDPHIPPVDP
		1				TTHLTFNHLSIVCVSLTLSLPHL
						GSPCLWSPALVSQEAASHQDRR
20880	51248	Α	20999	309	1310	RHQRAPEQGPDASWKMLDSLL
						ALGG\LVLL\RDSVE\WEG\RSLL
	ļ					K\ALVKKSALCGE\QVHILGCE\
						VSEKEFREGFDS**STIRLVYHD
						F\FRDP\LNWSKTEEGL/VPGGPL
						GALRS\MCKR\TDPVPVTIALDS
			1			LS\WLLLRLHCT\TLCQ\ALHAV
						EPSRTLVLGDSSSVGKVSVLGL
						LHERLPRPAPVVALSSLAQTKV
						TLGGYHGARPSAHIL\CRRPRQ\
						RPN\YQTQWFSI\LPDFSLGSPRG
		1				ALCRVPALIPIPHITPGGIPHNSF
		1				*PFNLHL\SKKEREARDSLILPFQ
l		1				FSSEKQQA\LLRPRPGQATSHIF
						YEPDAYDDLDQEDPDDDLNILP
	J	1	L	L	l	TEL DITTODD QUEDI DEDUCATED

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		,
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
20881	51249	A	21000	418	1304	GLVMEIGMHHHEHHHLQAPNK
						EDILKISEDERMELSKSFRVYCII
						LVKPKDVSLWAAVKETWTKH
	ļ					CDKAEFFSSENVKVFESINMDT
						NDMWLMMRKAYKYAFDKYR
		1				DQYNWFFLARPTTFAIIENLKY
		ŀ				FLLKKDPSQPFYLGHTIKSGDLE
İ						YVGMEGGIVLSVESMKRLNSL
						LNIPEKCPEQGGMIWKISEDKQ
						LAVCLKYAGVFAENAEDADGK
						DVFNSQICWAFY*R/WQ*LITPN
						QVVEGCCSDMAVTFNGLTPNQ
						MHVMMYGVYRLRAFGPYFSM
						MHWFSYLQMVLTMTEKW
20882	51250	A	21001	1	1092	
20883	51251	Α	21002	130	397	EGVHSQVNLFPIKGNQTHVPAS
		1			Ì	HGTDSALPPTLQMDYRALVHE
						PD*AAYGELRAMVLDLRAFYA
		ŀ				ELYHIISSNLEKIVNPKGEEKPF
20884	51252	Α	21003	11	805	SSMAKPC\GVRLSGKGRKQVEV
						FYRNLFQEAEEFLLR\FLPPESHI
						P*NRLLQEDFPFNVA*/LLNLSS
		l				GAPLDIPPSPDP\PP\KDDEMETD
		l				K\QEKKEVPKCDFLP\GNEKCPC
		l				PCLPLVKPRSLDLSKEKC\ILVI\
						TWGSRHLDPPRFEDGK*FWG*
		l				AIPGEGAWRG*NA\VKDPKWE
		l				AFPATPISQVLSSERGDACGPRP
}						PQETHV\MDYRALVHERDEAA
1		İ				YGELRAMVLDLRAFYAELYHII
						SSNL\EKIV\TPRGEEKPIYVLEP
20885	51253	A	21004	352	670	AKRSRIHVGVVIVDSDGAELGR
						DAAALSEKPPSLTIGLHMELVG
						MGGEPALTATPANCPEYPAKPS
						SI/EQQ*QSQVQ**Q*QVRHGSG
						PLHRTPRQLFLCSRQFAL
20886	51254	Α	21005	3	222	SPEESPPACPRQVFPLYHLQRG
		1				QGKRVQVG*GGSLWITTKTRLL
						PGGVGDNEVAPGSWHLALVFR
			<u> </u>			LLSSGRP

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20887	51255	A	21006		3639	MSEETLQSKLAAAKKKIQDILN VLVSDLNRSNGVVLPTLDKWK DAPEAWDLCLALRSICGLAIQL LGAAPLTRAANFSVAEVTDVR DSARRWETRKAPYLSCRIALRM NEAAVFRHLVPGVYFPRCDQG CQLTAALGCGAGVQSHVGPVT VTTGPDSVTAADAAGLCRVLR AVGWGGSECRLTPHPYKFHVE PYGGRLVPPRLSVDPGSSRREA TSKVAVDLLVLDAAGADGPVS QSPVWKEAAGVVSQLRGGRQ
20888	51256	Α	21011	137		KPRCCHSATMFEVPLVQGSILK KVLEALKDLINEACWDISSSGV NLQSMDS/SHVSLLQFTLPSEGF DTYRCDCNLAMGVNLTIMSKIL KCAGNEDIVTLRIEDNVNTLVL VFEASNQEKVSDYEIKLMDLD VEQLGIAEQEYHYVVKMLSGE FASIC*DLSHIGDAVVISGVKDG IIFSASGDLGN\KNKLSQTSTVD KEEEAVTMEMNEPVQLTFALR YLNFLTKATPLSSMVTLNMSA DIHLVVDYKIADVGHLKYYLA PKMEDEEGP
20889	51257	В	21012	205	1309	
20890	51258	Α	21013	189	410	
20891	51259	А	21014	1767	2158	RAQPHLYPEAARTRGRK*LHFC APSSAPRAPVSERR/CAGRFLSW CPVAAWDPARWARRARRRR AAARRRPPRWRRRCLSARGR RRSLNSSKTKNTKCLNSINQRL KILSLQKDLMCGTAGRCKTLTE Q
20892	51260	Α	21015	1	1710	
20893	51261	A	21016	132	522	
20894	51262	Α	21017	1	1386	
20895	51263	A	21018	331	3101	SFSEANISTAKKTFLYNELYVY NTRKDTWTKVDIPSPPPRRCAH QAVVVPQGGGQLWVFGGEFAS PNGEQFYHYKDLWVLHLATKT WEQVKSTGGPSGRSGHRMVA WKRQLILFGGFHESTRDYIYYN DVYAFNLDTFTWSKLSPSGTVP TPRSGCQMSVTPQGGIV\VYGG YSKQMPPHSGQSRGRKPGSCLL LLLLLLLLLLLLLCVLPLGYSL IELSSSYRLHGNEPVPATFHGLN SHMGLVATILDNVNP

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20896	51264	Α	21019	2	2911	VRGPRARCRRRSSKTEPRKGPS
						RTSTRKRLELGAGSDSISVRRAP
						PHEHLFPSGERGPFSFLVSRRGL
						GPGKMGKKGKKEKKGRGAEK
						TAAKMEKKVSKRSRKEEPGVC
						RRIRPDLGVECSAAAAHGFPNR
						GQFYGLPCARHSSLGFTPHEQE
		İ				DLEALIAHFQTLDAKRTQTVEV
		1				PCPPPSPRKCRGSQFVVTEDSVR
1						KGTALRLNASLSVHPEKDELIL
		İ				FGGEYFNGQKTFLYNELYVYNI
						RKDTWTKVDIPSP
20897	51265	Α	21020	1	208	DGYRHEMLPASLIQAQRDYFG
						AH/TYELWAKPGQFIH/TNWTG
						HGGTVSSSSYNAFALFSVQFFK
						SVVRDS
20898	51266	Α	21021	3	316	
20899	51267	Α	21022	2	578	LIGLAVMGQNLILNMNDHGFV\
		ļ				TVSKVDDFLANEAKGTKVVGA
						QSLKEMVSKLKKPRRIILLVKA
						GQAVDDFIEKLRRCRDLKAKGI
						LFVGSGVSGGEEGARYGPSLMP
						GGNKEAWPHIKTIFQGIAAKVG
						TGEPCCDWVGDEGAGHFVKM
						VHNGIEYGDMQLICEAYHLMK
						DVLGMAQDEMALCPIFCSVF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20900	51268	Α	21023	3	1758	AYANLGTRFWREPLRVFPSLVL
						RASPLFGSALSAAMAQADIALI
						V\LFVMGHNLILNMKDHGFEVC
						AFNRTVSKVDDFLANEAKGTK
						VVGAQSLKEMVSKLNKPPRDQ
						YLLC/VKAGQAVDDFIEKLVPL
						LDTGDIII\DGGNSEY*GTPQDG
						CRDLKAKGILFEGS\GVSAGEE
1						GD\RYGP\SLMPGGNK\EAWPHI
						KTIFQGIA\AKVGTGEPCC\DWV
					,	GDEGAGHFVKMVHNGIEYGD
						MQLICEAYHLMKDVLGMAQD
						EMAQAFEDWNKTELDSFLIEIT
						ANILKFQDTDGKHLLPKIQGTS
						AGARRGNRGSGT/WPFSGPGKY
						GVPV\TLI\GEAVF\ARCLSSL\KD
						ERI\QASKKLKGPQK\FQFDGD*
						EIISWEDIRKALYASKIISYAQGF
						MLLRHAA\TEFGWTLNYG\GIA
						LMWRGGCIIRSVFLGKIKDAFD
						RNP\ELQNLLLD\DFFKVQLFEN
						CQ\DSWR\RAVSTGVQGWAFP
						MPLFYRLPLSI*TGTSHEMLPAQ
						PSSRLQRGLLSGAHTL*NSWAK
ĺ						PGAVLFHTNWTG\HGGTVSSSS
					:	YQCLDHAAPVTLHDSTVPGHSI
						GLNGTAHLALCPIFCSVFFKSV
20901	51269	Α	21024	317	885	MRWRSKSAWRAEPKSRTNSGT
						ASNVSPSRPLEC*R/FLRLVSGE
						LILALIAFICIETIMACSPCEGLY
ŀ						FFEFVSCSAFVVTGVLLIMFSLN
						LHMRIPQINWNLTDLVNTGLSA
						FLFFIASIVLAALNHRAGAEIAA
						VIFGFLATAAYAVNTFLAVQK
						WRVSVRQQSTNDYIRARTESRD
						VDSRPEIQRLDT
20902	51270	Α	21025	4761	5127	LGSGDLPWEINPLSSCSLLREKD
	ļ	İ				PPTTSGPQT\TSPRNISPISNPHTR
						TSKRLNRSGQAFLQNLLPQELA
						TSARNLTTRPRNACSPGFLLSR
						VPSVRDPTGNWTVQLTWQPLS
						EPLELWPKAL

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20903	51271	A	21026	174	2238	LRSGDLPWEINPLSSCSLLHEKD PPTTSGPQT\TSPRNISPISNPRQR RHILSMDPKLRRRSWTWVGSLP LVFNHCRDASLIIHPGFRGVRPR RDVCLGPSPLATITDAELYVTL TVEGKSIPFLIDTEATHSTLPSFK GPVALASIHVQNVQIKRDKEGH YIMVKGSIQQEELTILNMYAPN TGAPRFIKQVLNDLQRDLDSHT IIMGDFNTPLSTLDRSTRQKVN KDIQDLNSALHQADLIDIYRTL HTKSTECTFFSAPHRTYSKIDHI VGSKALLSKCKRTEIIINCLSDH SAIKLELRIKKLTQNHSTTWKL NNLLNDYWVHKEMKAEIKM FFEINENKNTTYQNLWDTFKAV CRGKFIALNAYKRKQERSKIDT LTSQLEELEKQEQTRSKASRRE EITKIRAELKEIETQKTLQKINES RSWFFEKINKIDILLARLIKKKR EKNQIDAIKNDKGDIATDPTEIQ TTIREYYKHLCPNKLENLEEMD KFLDTYTLPRLNQEEDESLNRPI TGSEIVAIINSLPTKKSPGPDGFT AKFYQRYKEELVPFLQKYKEEL VPFLLKLFQSIEKEGILPNSFSEA SIILIPKPGRDTTKKENFRPISLM NIDAKILNKILANQIQQHIKKLI
						HHDQVGFIPGMQGWFNIRKSIN VIQHINRTKDKNHMIISIDAEKA FDKIQQPSC
20904	51272	A	21027	1 .	181	MEEKEERKRRKRKRREEGR DGERKRMEKQKENQDPKDPPT TSGPQT\TSPRNISPISNP
20905	51273	A	21028	167	408	LGSRTFPWEINPPSSCSLLREKD PPTTSGPQT\TRPRNISPISNLVS GLFLLSSPTSLTIPQPLSSFNLDD TLQSLPSLNF
20906	51274	A	21029	449	663	LGSGDLPWEINTQSSCSLLCEK DPPTTSGPQT\TSPRNISPISNPDP TGNRTVQLTWQPVPEPLELWP KAL
20907	51275	A	21030	79	381	LGSGDLPWEINPLSSCSLLREKE PPTTSGPQT\TRPRNISPISNPELA TLAGNLATGPRNARSPGFLLSH VLSVWDPTENQTVQLTWQPLP QPLELWPKAL
20908	51276	Α	21031	122	334	LRSGDLLWEINPLSS*SLLHEKD PPTTSSPQT\TSPRNISPISNPDPT ENQTVQLTWQPLPEPLELWPK AL

SEQ ID			SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20909	51277	A	21032	109	320	LGSGDLP*EINPPSSCSLLREKD PPTTSGPQT\TSPRNISPISNPDPT GNRTVQLTWQPLPQPLELWPK AL
20910	51278	A	21033	620	717	
20911	51279	A	21034	3	678	GGGFSDRDRTIALQPGQQIWM WGIGKQVGGCGWLRGKNRVV GAAKDQSGTGEKFYFIPKAMG AMEGSGQRRTLPDPGAHRNPS GGYSATDCEGSRPAPGEINSHA SHTKPVWWFFTQTRMKFGAVT RIGGLPWEVNPLSSCSLLRE/KD PPTTSGPQSNQPRKHPPISNPEL VHVPEIWPLGQGMPAARIPPKP RPICVGPH*KSTVQLTWQPLPEP LELWPRLSD
20912	51280	A	21035	339	1098	LGSGDLPWEINPLSSCSLLREKD PPTTSGPPT\TSPRNISPISNPGET KETRFIRGPKTPAPVTDWEGSL PLVFNHCRDASLIIHPRFKGVRP RRDACLGPSPLAASPTFLGKGQ RLKTDTARLPRKPPRPSRTLSFS TAPRILSIVVPGSGHTLAFLDVT RHRLYRESKPVVGEEVDYSEW LSDVGKSEWSFSTCASSTSVNK HDTGRRLLGKRGGGMECSKGL CNFITMTKNIRSLAQMTIWQYL SPCREPV
20913	51281	С	21036	1	438	
20914	51282	A	21037	174	483	LRSGDLPWEINPLSSCSLLHEKD PPTTSGPQT\TSPRNISPISNPRQR RHILSMDPKLRRRSWTWVGSLP LVFNHCRDASLIIHPGFRGVRPR RDVCLGPSPLA
20915	51283	A	21038	590	824	PNSSWMRGEPKKDPPTTSGPQT \TSPRNISPISNPGRLSDYSPRFQ RCQTTQGRLPWSFTLSGKSRFS GEGATWGLKT
20916	51284	В	21039	1	1596	
20917	51285	В	21040	155	2206	
20918	51286	В	21041	1	831	
20919	51287	В	21042	75	474	
20920	51288	В	21043	38	1009	
20921	51289	С	21044	335	481	
20922	51290	В	21045	66	652	
20923	51291	В	21046	1	673	
20924	51292	В	21047	1	451	
20925	51293	В	21048	74	3516	
20926	51294	В	21049	783	3463	
20927	51295	В	21050	1	807	
20928	51296	В	21051	41	1700	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
20929	51297	C	21052	18	381	
20930	51298	В	21053	1	3147	
20931	51299	В	21054	1	2586	
20932	51300	С	21055	756	1403	
20933	51301	В	21056	42	205	
20934	51302	В	21057	40	420	
20935	51303	В	21058	303	1170	
20936	51304	В	21059	1	2631	
20937	51305	В	21060	19	3093	
20938	51306	В	21061	23	241	
20939	51307	В	21062	84	1775	
20940	51308	В	21063	68	1542	
20941	51309	В	21064	74	2566	
20942	51310	С	21065	730	1023	
20943	51311	В	21066	67	1159	
20944	51312	В	21067	294	929	
20945	51313	В	21070	68	336	
20946	51314	В	21071	1	1992	
20947	51315	В	21072	1	4188	
20948	51316	С	21073	44	535	
20949	51317	В	21074	983	2636	
20950	51318	В	21076	289	1693	
20951	51319	В	21077	364	2393	
20952	51320	В	21078	471	590	
20953	51321	В	21079	1	1026	
20954	51322	В	21080	343	4253	
20955	51323	В	21081	1	819	
20956	51324	В	21082	648	894	
20957	51325	В	21083	1	1236	
20958	51326	В	21084	Ī	822	
20959	51327	В	21085	211	363	
20960	51328	В	21086	193	264	
20961	51329	В	21087	1	1260	
20962	51330	В	21088	143	3158	
20963	51331	В	21089	85	353	
20964	51332	В	21090	221	598	
20965	51333	В	21091	1	4878	
20966	51334	В	21092	171	362	
20967	51335	С	21093	320	408	
20968	51336	В	21094	1	1218	
20969	51337	В	21095	1	1761	
20970	51338	В	21096	315	1224	
20971	51339	В	21097	526	3482	
20972	51340	В	21098	41	4858	
20973	51341	С	21099	38	250	
20974	51342	С	21100	157	225	
20975	51343	С	21101	191	415	
20976	51344	В	21102	42	507	
20977	51345	С	21103	71	367	
20978	51346	С	21104	231	439	
20979	51347	С	21105	207	459	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20980	51348	Α	21106	11	429	
20981	51349	Α	21107	184	544	VRLLKILQLQRFKMYVQQLDL
						TYFLRKIKCTLENGIVMSNTEA
						ESGVQLKPGRWGASALYSSHHI
						IH*A*LLVPR*VSNVVCSRNDT*
						TKNKDTKNRRC*PKSSTRRGK*
			:			KRERKEKEVT
20982	51350	Α	21108	3	1200	LSQKEEEEDTFIEEQQLEEEKLL
						ERERQRLHEEWLLREQKAQEE
						FRIKKEKEEAAKKRQEEQERKL
						KEQWEEQQRKEREEEEQKRQE
						KKEKEKTFMWKPSVDLFSHLA
						SLRTPIICQDKGNAYQEHFPEK
						KKKRSSQLAIPEGAFGYPHHLL
						AILGLRCSRKHNFPTSSPTLLIKS
						MFTTFGMEQCRRDDYDPDASL
						EYSEETYQQFLDFYEDVLPEF
			,			KNVGKVIQF/KGLFEIQQCPRGK
						HCNFLHVFRNPNNEFWEANRDI
						YLSPDRTGSSFGKNSERRERMG
						HHDDYYSRLRGRRNPSPDHSY
						KRNGESERKSSRHRGKKSHKRT
						SKSRERHNSRSRGRNRDRSRDR
						SRGRGSRSRSRSRSRRSRSRSQ SSSRSRSRGRRRSGNRDRTVQS
20983	51351	A	21109	789	907	SSSKSKSKUKKKSUINDKIVQS
20984	51352	A	211109	236	1823	CPIQRQSPGPAQTGRWEPLPCT
20704	31332	^	21110	230	1023	VPITLRMSEGSGGAGKMAALE
						KMTFPKKMTFPEKPSHKKYRA
						ALKKEKRKIRRQELARLRDSGL
		1				SQKEEEEDTFIEEQQLEEEKLLE
						RERERLHEEWLLREQKAQEEFR
	1		;			IKKEKEEAAKKRQEEQERKLKE
						QWEEQQRKEREEEEQKRQEKK
						EKEEALQKMLDQAENELENGT
						TWQNPEPPL\DFRVMEKDGVN
				·		CPFYSKTGACRFGDRCSRKHYF
						PTS\SPTLLIKSMFTTLGMEQC\R
						RDDYDPDASLEYSEEETYQQFL
						DFYEDVLPEFKNVGKVIQFKVS
						CNLEPHLRG\NIYVQYQSEEEC
		İ				QAALSLFNGRWYV\AGRQLQC
						EF*TGDRWKMAICGLFEIQQCP
						RGKHCNFLHVFRNPNNEFWEA
						NRDIYLSPDRTGSSFGKNSERRE
						RMGHHDDYYSRLRGRRNPSPD
						HSYKRNGESERKSSRHRGKKSH
						KRTSKSRERHNSRSRGRNRDRS
1						RDRSRGRGSRSRSRSRSRRSRRS
L						RSQSSSRSRSRGRRRSGNRDRT
20985	51353	Α	21111	289	366	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20986	51354	Α	21112	1	1289	
20987	51355	A	21113	60	367	EELGYAGRGMGARQPPPARKA VSPLQERAEPVGSPG*GWRSRP STHGIASHGGPACSLCSLHFTFF VDFPSLFSPICLPEGWDWQQEQ PCLLYLSSPRCTQA
20988	51356	A	21114		653	RRRGSAAGGRAAERKAEVGA MNMAGGKAGKDSGKAKAKA V\SRSQR\AGLQFPVGRIHRHLK TRT\TSHGR\VGATAAVY\SAAI L\EYLT\AEVLEL\AGYAS\KDLQ SKSVSLPRHLQLAIPWVIEELDS LIQGLP*AGGGVIPSHPHKS\LIG KEGDSKKTALEGMLLTQPLSLP RHLYCNWDKKNNGDMWNFLP PVKWKSIDNYCRHDKRNICMF LDSEV
20989	51357	Α	21115		1019	SVCWAGGAPRAHSTRTHAWLG RPAMEPGSVENLSIVYRSRDFL VVNKHWDVRIDSKAWRETGRP EP\QVASVEESGSDSGLC/HHSR FCHQLDFSTSGALCVALLVLGG LPGSTGLPRELRAILPLLQLRGH IQESRVSIVRWGGPGPRVSPTNL TPAFCAPGCENPKPSLTGSWSG R*VWIGPAPSLIRHRPHPAGRTH QLRV\HCSALGHPVVGDLTYGE VSGREDRPFRMMLHAFYLRIPT DTECVEVCTPDPFLPSL\DACWS PHTLLQSL\DQL\VAGLKGQPPD P*TRG*GGPRPGSPSALLPGHGR PPPPTTKPPETEAQRGPCLQWL SEWTLEPDS
20990	51358	Α	21116	698	1472	RWAPRRGLSVDLDGGSSAPLCP PATTARLDAEPHGSLPNSCLWE CSCLLQGIWAAGPSGLSLHYPP LSSPLGSSPPSRPQRLVLIRAGR GGWVGSVGETMTSSYGHVLER QPALGGRLDSPGNLDTLQAKK NFSVSHLLDLEEAGDMVAAQA DENVGE\SSRTRGGGAGLGKGL SGVQTQRASAVWESSRLKMLS RVGRASLGQRMTA*GRLQLYP HAPKSVTDRSPCSSQVTKSWNR GALQHSMIFTKRCLFQTLKE

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	1	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
20991	51359	A	21117	50	1226	KMAEEVVVVAKFDYVAQQEQ ELDIKKNERLWLLDDSKSWWR VRNSMNKTGF\VPSTYVGR\EN SARKASIVKNLKDTLGIGKVKR KPSVPDSASPADDSFV\DPGERL Y\DLNMPA\YVKFN\YMAERED ELS\LIKG\TKVIVMEKCS\DGW WRGTYNGQVGWFPSNYVTEE\ GESPLGDHVGSLSEKLAAVVN NLNTGQVLHVVQALYPFSSSN DEELNFEK\GDVMDVIEKPEND PEWWKCRKINGMVGLVPQKL W*PLLQSIQLTSGLGTLTFQCGF IRPYIPGKFVGNPWYYGKVTRH QAEMALNERGHEGDFLIRDSES SPNDFSVSLKAQGKNKHFKVQ LK\ETVYCIWQRKSCTME*LVE HYK\KAPNFYNVNKGEKLYLV
20992	51360	Α	21118	132	428	
20993	51361	A	21119	438	1066	AKLAGAAVPVCARGRAAPLAA AALPAARL\PSGAGPAGLVSDV RRRPFARE/PEPRVP/SQPPALSA LGALALKLFSSPPSCWSFSVKT GVLVPWIMCISWPYFLSGNLLT MMW LNQKGKGGRGEALKNDLVVAP PPSPRRGVR*EALHVPDTGPFK
						KPRKRK*NQRQREAERK*TFPR EEEARRETHGPLPAPRPPPPLCP/ PAASSAPRPGPKGAR*LSLSAE GRARRRSAGASGGAGAAATAA AAAAPEHPRGSEPRQPASPAPQ RERPRRPAPRLPPQCPRPSKGRA WTLQTTTGQKVCTGAGTRRA WRRAGAVRRAFWA
20995	51363	A	21121	1	924	
20996	51364	Α	21122	1	435	
20997	51365	A	21123 21124	1	524 602	MGKGSPAHAQGPGSFLHGGAG DQGRGRPITLYEDRGFQG\RHY ECSSDHPNLQAYLSRCNSVRVD SGCWVLYEQPNYSGLQ*FLRR GDYADHQQWMGLSDSVRSCR LIP\HASSHRLRIYEREDYRGQM VEITEDCSSLHDRFHLSEIHSFN VLEGSWVLYELPNYQGRQYLL RPGDCRWCQDWGATDARVGS LRRAVELY
20999	51367	Α	21125	1	1239	
21000	51368	A	21126	298	408	
21001	51369	С	21127	92	241	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21002	51370	A	21128	33	191	RDDPRVRPPPNSHT*PQQEPGL* LIKCTSPPQAPAPRTVHGPYFY MRLIKMF
21003	51371	Α	21129	3	1285	
21004	51372	A	21130	3	3468	QPGHTIYLLPTVVICNLLPCELD FYVKGMPINGTLKPGKEAALH TADTSQNIELGVSLENFPLCKEL LIPPGTQNYMVRMRLYDVNRR QLNLTIRIVCRAEGSLKIFISAPY WLINKTGLPLIFRQDNAKTDAA GQFEEHELARSLSPLLFCYADK EQPNLCTMRIGRGIHPEGMPGW CQGFSLDGGSGVRALKVIQQG NRPGLIYNIGIDVKKGRGRYIDT CMVIFAPRYLLDNKSSHKLAFA
21005	51373	A	21131	3	451	QREFARGQGTA
21005	51374	A	21132	2	107	LEGLEES/VRDRCKKEFTDLMIE MDSRLLGPTSKVK
21007	51375	A	21133	2	158	PGIPVL/DYRTYTDGFFSLPSKD GDKDVMITGKLDILKPQNQHR VFTVESRVI
21008	51376	A	21134	2	5632	FVRRTCRESRCWPGAETELDH WSLGEGVRGAMALQLWALTL LGLLGAGASLRPRKLDFFRSEK ELNHLAVDEASGVVYLGAVNA LYQLDAKLQLEQQVATGPALD NKKCTPPIEAS\QCHEAEMTDN VNPLLLVDPPRKRLVECGQLLK GI\CALRALSNISLRLFYEDGSG EKSFVASNDEGVATVGLVSSTG PGGDRVLFVGKGNGPHDNGIIV STRLLDRTDSREAFEAYTDHAT YKAGYLSTNTQQFVAAF
21009	51377	A	21135	168	5568	AREKCGGAMGAGSSTEQRSPE QPPEGSSTPAEPEPSGGGPSAEA APDTTADPAIAASDPATKLLQK NGQLSTINGVAEQDELSLQEGD LNGQKGALNGQGALNSQEEEE VIVRQVDHIDSEDVSDRDSDTE MTTKSAVVHDITDDGQEETPDI IEQIPSSESNLEELTQPTESQAND IGFKKVFKFVGFKFTVKKDKTE KPDTVQLLTVKKDEGEGAAGA GDHQDPSLGAGEAASKESEPK QSTEKPEETLKREQ

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21010	51378	A	21136	807	1162	ISYHGGRRRRPQ/PVGPLGSCRS
						*SSRRWSTTGRTTFRAGWKSCS
		1				TPPSTSSLPTSTGTW*GPGTSAL
						LPPTPLPPPRAAATPCACACASE
						SRMPAARALGLRVRRRLLEGRS
				:		GKDTQG
21011	51379	A	21137	1	562	
21012	51380	A	21138	1	423	
21013	51381	A	21139	1	1871	MTTFKEAMTFKDVAVVFTEEE
						LGLLDLAQRKLYRDVMLENFR
						NLLSVGHQAFHRDTFHFLREEK
						IWMMKTAIQREGNSGDKIQTE
						METVSEAGTHQEWSFQQIWEKI
						ASDLTRSQDLVINSSQFSKEGDF
Ì			1			PCQTE/ARTICNSHKTEI/FPRA/I
						GYKPSFSDVSHFDFHQQLHSGE
						KSHTCDECGKNFCYISALRIHQ
						RVHMGEKCYKCDVCGKEFSQS
						SHLQTHQRVHTGEKPFKCVEC
}						GKGFSRRSALNVHH
21014	51382	A	21140	1056	2988	FCILIGDKIQTEMETVSEAGTHQ
		1				EWSFQQIWEKIASDLTRSQDLV
						INSSQFSKEGDFPCQTEARTICN
						SHKTEIFPRAIGYKPSFSDVSLF
						DFHQQLHSGEKSHTCDECGKN
		ı				FCYISALHI\HQRVHMGEKCYK
		ı				CDVCGKEFSQSSHLQTHQRVH
						TVEKPFKCVECGKGFSRRSALN
1						VHHKLHTGEKPYNCEECGKAFI
				•		HDSQLQEHQRIHTGEKPFKCDI
		ı				CGKSFCGRSRLNRHSMVHTAE
						KPFRCDTCDKSFRQRSALNSHR
						MIHTGEKPYKCEECGKGFICRR
						DLYTHHMVHTGEKPYNCKECG
						KSFRWASCLLKTSSESHSG\EKP
						FKC\EECGKGFYTISQCYS\HQR
		ŀ				SHSGEKPYKCVECGKGYKRRL
	1					DLDFHQRVHTGEKLYNCKECG
			:			KSFSRAPCLLKHERLHSGEKPF
						QCEECGKRFTQNSHLHSHQR\V
						HTGEKPYKCEKCGKGYNSKFIL
						YMHQ\RVHTGERPYNCKECGK
		1				SFGWASCLLKHQRLHSGENPFK
		1				CEECGKRFTQNSQLHSHQRVH
						TGEKPYKCDECGKGFSWSST\R
						LTHQR\RH\SRETPLKCEQHG\K
						NIV\QNSFS/EKLPEKVHQC*KK
						PYKCEDL/CGKGYNRRL*IL\DM
					-	HQR\VHMGEKTWKCRECDMCF
						\SQAS\SLRLHQNVHVGEKP
			<u> </u>	J	1	

SEQ ID			SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide sequence	hođ	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	Sequence		05/210,217	sequence		
21015	51383	A	21141	154	11470	  KDGDHMGYTYKVIERILEIGAF
21013	31383	A	21141	134	1470	RPHMSLGQLQSARGVACLGCK
		1				GTCSGFEPHSWRKICKSC\NAA
ļ						KRTTA*HLT*KNDRKLGRWLR
ŀ						DSKYSTLHLLG*KAGTASGFYK
İ						
						RNP\MI\LTNPIATGKDPTFDTIT
						YEWGSPWEFTQETGDCSYMGA
						HSPREEGSQLTGHQRGAFYRA
		ŀ				ARPAHAPGLPIYD\QDPLALPVD
						FWRNELKTEWKEFVK\KYKSE
						GPSALGEVALPGQGWLAPRKE
						GE/GSRKKPEGGRRP/PAATTNG
		ļ				SLSDPSKEVEYVCELCKGAAPP
						DSPVVY\SDRAGYNKQWHPTCF
						VCAKCSEPLV\DLIYFWKDGAP
						WCGRHYCESLRPR/CAPACDEI\
						IFAEDYQRVEDLAWHRKHFVC
						EGCEQLLSGLGVHRHPGVQLL
						VPQLAASPKRSLKGCPTPQPESP
						GIPTEKDKPGCARDPSLRVPDV
						TSKPSEINQ
21016	51384	Α	21142	1	1332	
21017	51385	C	21143	147	272	
21018	51386	Α	21144	160	374	ETPVSSRCEKHVSSPHLCPGIHR
						AVSLSSH\STCGRSWRSELHGPQ
						STNLSGNTELPLHFLFGAWQTG
						VFKM
21019	51387	Α	21145	3	4366	LGAQQRELQEALGARAALEAL
						LGRLQAERRGLDAAHERDVRE
						LRARAASLTMHFRARATGPAA
						PPPRLREVHDSYALLVAESWRE
1						TVQLYEDEVRELEEALRRGQES
						RLQAEEETRLCAQEAEALRREA
						LGLEQLRARLEDALLRMREEY
						GIQAEERQRAIDCLEDEKATLT
						LAMADWLRDYQDLLQVKTGL
						SLEVATYRALLEGESNPEIVIW
						AEHVENMPSEFRNKYYHYTDS
					Ì	LLQRENEWNLFSRQKAPLAS
21020	51388	Α	21146	1	4617	RVIDCLEDEKATLTLAMADWL
ŀ			1		1	RDYQDLLQVKTGLSLEVATYRI
					1	VEHVAVEGVSKLWKPLWQKS
						QSLVISSGVSFTGGRLNLERLRR
						GPTPLASTESVKQVLVNNRMII
					1	DDEMVKTLAMSRTCSAPNPLT
						ADVKTWSRSYRSEDAATGPVL
						GVASTVALLEGESNPEIVIWAE
					1	HVENMPSEFRNKSYHYTDSLL
[					[	QRE\NERNLFS\RQKAPLASFNH
						SSALYSNLSGHRGSQTGTSIGG
						DARRGFLGSGYSSSATT
L	L			<u> </u>		DARKOI EGOG I GOOM I I

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21021	51389	Α	21147	2	435	
21022	51390	Α	21148	1	1233	MCEEETTALVCDNGSGLCKAG
						FAGDDAPRAVFPSIVGRPRHQG
						VMVGMGQKDSYVGDEAQSKR
						GILTLKYPIEHGIITNWDDMEK
						KKEAEELARNMAIQQELEKEK
						QRVAQQKQQQLEQEQFHAFEE
						MIRNQELEKERLKIVQEFGKVD
						PGLGGPLVPD/SGEALLRCVPHL
į						NSLIHTAFRLSHNCKAS*ATCG
						GQVLETWSTEQLRKYSHNRWI
						APCGGAWAAVPTVSPVSQCQH
						CPGSGDMWNSLWKTDEE*IYH
İ						YPCSHPQAKCWV*LLQHRERR
			1			RTFPHTGSAGPHHTGLDSCKQF
		1				*AVREHGVR\GIHETTYNSIMKC
İ		İ				DIDIRKDLYANNVLSGGTTMYP
						GIADRMQKEITALAPSTMKIKII
•		ļ				APPERKYSVWIGGSILASLSTFQ
						QMWISKPEYDEAGPSIVHRKCF
21023	51391	Α	21149	2	1273	PRVRPRVRPRVRSLGEEPLIPSV
		1				LRSPAHSATHTMCEEETTALVC
		l				DNGSGLCKAGFAGDDAPRAVF
		1				PSIVGRPRHQGVMVGMGQKDS
		1				YVGDEAQSKRGILTLKYPI\EHG
		l				\IITNWDDMEKIWAPLLLTMEL
		ļ				RVSTWKRHPTLLTEAPLNPKAN
		ł			į	REKMTQIMFETFNVPAMYVAI
İ		i				QAVLSLYASGRTTGIVLDSG\D
Ì						GVTHNVPIYEGYALPHAIMRLD
					1	LAG\RDLTDYLMKIL/AKRGGY
		ł				SFVTTAEREIVRDIKEKLCYVAL
						DFEK*\MATAASSSSLE\KSYELP
		1				DGAGLSPIGKWTLP\CPETLFQP
		1				YFISNGVPLEFNDDNLTIPSMK\
		1				CDIDIP*GLICPTMFLSGGTTHV
		]	1			PWALLDRIQKENPSPGPPGTMK
		}				IKI\IAPPEREVTSVWD/LGGSILA
		1				SLSTFQA\MWISKPEYDDAGPSI
				<u> </u>		VHRKCF
21024	51392	Α	21150	95	674	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21025	51393	Α	21151	1	1536	MGGGEQGTLRTAENQGLQVCP
121023	3.373	ļ .	21101	•		GEAGSSLGFSGNTPTNGLPARG
						DQTSYRDFWSGPQFPDTSLRTA
						APGAREQCGTHGGEGSTLEGSR
						PDVPQDLFSPTLGFYGSTYPELE
						QGMLGGH*G/PAENQGLQVCP
						GEAGSSLGFSGNTPTNGLPARG
						DQTSYRDFWSGPQFPDTSLRTA
						APGAREQCGTHGGEGSTLEGSR
						PDVPQDLFSPTLGFYGSTYPELE
						QGMLGGHVEASREYQMLSHV
						GNPDYPHALATEQATSPSTESG
						SAGFKGVALAMLSKITMCVRIG
						ILVFQQDQGSMCKAFSTTGMG
		-				LVTYLLTCRAWSRLQCGASYR
						AHDVNVLPGCSGPAALLLGDW
						FSLHSSDAVLPAQGQGMLSQG
						EADACITLPHLFPSREVVFLLGL
						EAGGLIPGAEGPPEDMGLVRGP
						PAVAAGPPPRWLSPEFCLLQYS
						EEEGCKEPLWLKVEAEVSTLYR
						FMPTGFTRGPFNIEHREWGNTT
						ATWTRMISNHNKHSQRGRKMT
						ARLPCGGAPHFSDRTDDKWRR
						SSLLTRSGCQTERLLTSQTEPLP
		1				AGGSPHFSNGETGQRRSSPPQR
						GCDRPQALLTSKTEQQRKSASH
						ISYDERPGKDAPHFLDGMVAG
						KRRSSLSRLGSQA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
1	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21026	51394	A	21152	689	2176	PATSPTSRAMAQPSAPAASMLL
1020	31371	ļ.,	21132	00)		PCPECPSGPRFMTCCLSVSTFCS
						PSLRRPPGSIKGSSGLGGG\SSRT
						SCRLSG\ALGAGSCRLGSAGGL
						GSTLGG\NSYSSCYSFGSGGGY
						GSSFGG\VDGLLA\GGEKA\TMH
						NLIDF\LASYLDKVPCPWKEAN
						TELEV\KIRD\WYQRQ\APGPAR
						DYSQYYRTIEELQNKILTATVD
						NANILLQIDNARLAC/ADDFRT
						K\F\ETEQALRL\SVEADINGL\R
]						RVL\DELTLARADL\EMQIENLK
						EEAWPTLKERTTEGRRLNALR
						GQVGW*RSMLEMDAAPGVDL
						SRILNEMPSCVDHRVTRCINRP
						GAELKLGSGGWTEELKREVAT
					:	NSELVQSGKSKISEL\RRTMQAL
						EIELQSQLSMKA\SLEGNLGGD
		l				RATAYCVQLS\QIQGL\IGSGED
		l				QL\AQLR\C\EMEQQ\NQDYKILL
		}				DVKTRLEQEIATYRRLLEG\ED
		1				AHLT\QCKKEPVTTRQV\RTIVG
		1				KRSRD\GKVILLPRSRFHQTTR
21027	51395	С	21153	58	252	
21028	51396	Α	21154	2	190	
21029	51397	Α	21155	2	432	KMADGVDHIDIYADVGEEFNQ
		l				EAEYGGHDQIDLYDDVISPSAN
						NGDAPEDRDYMDTPPPVPGYG
						PPPGPPPPQQGPPPPPGPFPPRPP
		ŀ				GPLGPPLD/TLAPPPHLPG\PP\PG
						\APPPA\PHVEPSFLFLPPINSGH
L						GLHSDRPRFH
21030	51398	A	21156	1	1070	MGGIGARAAAAGKGRRRRARA
		ŀ				PPGLPPPTSDGGGGAAAASSLR
		ľ				RLHAAPPPGPSAAGSASRVPPTI
		ľ				GSEAAPAREDRRQLHSGRVRAS
						GSPRLTRPRPVKRRGRTRGGGA
						PRQGSAARPRGGKGAHSAVAS
						ARAAAGPGSASRPHGAASAAG
						TWALTYLVELFADIRVNVYVV
						HAVRHLPSACAAAAAAASCRF
						ALVGVGSEASSKKLMDLLPKR
						ELHGQNPVVTPCNKQFLSQFE
1						MQSRKTTQSGQMSGEGKAGPP
						GGSSRAAFPQGGRGRGRFPGA
						VPGGDRFPGPAGPGGPPPPFQD
						PSVNSDPVCGTFG/EH*PPGYQT
						PSNLSVRPMAKPKPEGTDLSRS
						REGFFKSRFAPN
			I	L	·	<u> </u>

SEQ ID	SEQ ID NO:		SEQ ID NO:	Nucleotide		Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21031	51399	A	21157	63	1750	MKMAGGVNHIDIYADVGE*FN
		İ				QEA\EYGGHDQIDVHDDAISPS
						ANNGDAPEDRDYMDTLPPTVG
						DDVGKGAAPNVVYTYTGKRIA
						LYIGNLTWWTTDEDLTEAVHS
:						LGVNDILE\IKFFENRANGQSKG
						FALVGVGSEASSKKLMDLLPK
						RELHGQNPVVTPCNKQFLSQFE
						MQSRKTTQSGQMSGEGKAGPP
						GGSSRAAFPQGGRGRGRFPGA
		ŀ				VPGGDRFPGPAGPGGPPPPFPA
		ŀ				GQTPPRPPLGSFQGPPG\PPGPPP
						PGQVLPPPLAGPPNRGDRPPPPV
						LFPGQPFGQPPLGPLPPGPPPPV
						PGYGPPPGPPPPQQGPPPPPGPF
						P\PRPTGSNLGHPLTLAPPSASFP
						GP\PP\GAPPPSPHVNPAFFPPPT
						NSGMPTSDSRGPPPTDPYGRPPP
						YDRGDYGPPGREMDTARTPLS
						EAEFEEIMNRNRAISSSAISRAV
				:		SDASA/GVDYGSAIETLVTAISLI
						KTIPKYPADD/RCTKFLISSLQD
]						CLHGIESKSYGSGSRRRERSRER
1		l				DHSRSREKSRRHKSRSRDRHDD
						YYRERSRERER\HRDRDRD\RD
21032	51400	A	21159	1	96	RERDREREYRHR TNFKSGLLGLAAED*HCPIASE
21032	31400	A	21139	1	90	AP*TITDAEL
21033	51401	A	21160	1	1056	MWKGPKGLDMYGKSSVSPKTS
21033	31401	^	21100	1	10.50	
						DILGRDTLLLALKVQTVVLQTA CGEGHVAGNCGRPLETEGSLQ
						`
		1				LTATKKLRDSVLQPKSPEFCQQ
						FTRAWNRTQVPDETEAPAGTY
						AAQSGDLPWEINPLSSCSLLHE
						KDPPTASGPQTDQPKEHLTNFK
						SGFRGVRP/LQGRLPWSFTLSGK
						SRFSGEGASTPTPSLCVSTPSPPF
						WGARNTQPLLLHPEWQ/CPLF*
						RSKYPNLVSLCPSPLFPRPDLLS
1						L/SAQSLISAPQPLISLCPNPLFPC
						PDLLSLHPNPLFPRPNPFPAFLE
						GACYKCQKSGHQAKECLQPGI
						PPKLRPICAGPHWKSDCPTHLA
						ATPRAPGTLAQGSLTASQIFLA
21034	51402	С	21161	283	408	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion; \=possible nucleotide insertion)
				sequence		
21035	51403	A	21162	3	505	ILNPVSGLFLFSSPLSLNLSSLSL
						NLFLLSILAPHFNLSILLISIPFIFP
	i					LEIELSNSPGSHSQSPWNSGPRL
						SDSFPDLLGLAAEE*HCPIASEA
						FSQTYRAYLSLSLSLSLSPEPPT
						AGDGVT/HSSPVASTTITVLGQT
		ļ				/SKPAQCWPLPKVCCHHSLTSD
						YVHPRP
21036	51404	Α	21163	164	420	LANFCEESGAGEPGSHSQSPWN
						SGPRLSDSFPDLLGLAAEDRHC
						PIASEAPLLQRKSLYKSPCTSSTI
						S*CCIPRSGISESLSVN
21037	51405	Α	21164	50	298	
21038	51406	Α	21165	49	433	SSSGIWPSHSQSPLRSSPAKLLS
			:			RFRCSLSSLLLCLATLLLVWQP
						LPEPLELWPKA/HLTDSFPDLLG
						LAAED/W/HCPIPSEAP*TLTDAE
						LRVTLTVEGKSIPFGAGKEVTL
						ESYWVSLTLSFLICYMII
21039	51407	Α	21166	282	374	CDLFLPHPN*SMYFVISPTPKKV
						LCNSPHT
21040	51408	Α	21167	71	333	LRSGDLPWEINPLSSCSLFREED
1						PPTTSGPQTNQPKEHLTNFKSA
	Ì					AED*RCQFASEAP*TITDAKLRV
						SLTAEGLLATLFLNPKPEY
21041	51409	Α	21168	113	342	AFGCFTTGVGRVTDSDGPLYTG
						PWAPGSHSQSPWNSGPRLSDSF
	:					PDLLGLAAED*HCPIASEAPFLT
						LIFLLCVIQ
21042	51410	A	21169	301	608	TFTPLKSYIKAPSSAGPVAEQE
	1					MTPEGKVGAGPE*LTWKNQFL
						FY
21043	51411	Α	21170	2	724	FLVSLMLPPAVVMLNHCYSLIF
						TNALVKRLFAWNMLSELATCA
						GNLATGPRNARSPGFLLSCVPS
						VRDPTGNQTVQLTWQPLPEPLE
						LWPKA/HLTDSFPDLLGLATED
						*HSPIASEAPWAIMDAELRVTL
						TVEDRDSSSSSSSSSSSSSSSSS
						SSSSSSSSSSSSSQSSSSSSSSPCD
						LFLPHPN*SMYFVISPTLKKVLY
						NVTHP*EVSL*FSPPLRMYFVRS
						IPCPQNIALNSTAYP
21044	51412	Α	21171	442	501	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
21045	51413	I A	21172	266	1003	SCMCLALDPCNHGEGHQPDFS
						GVKFRRHGADNHEASAATATT
						AAATTVAAAAAAAAAAAAAA
						VTLTGSHFQSPWNSGPRLSDSF
						PDLLSLATED*RCPIASEVPWTI
						TEAELRVTLTVEGKSIPCLIDTG
						ATHSTLPSFQGPVSLAPITVVG\I
						DGQASKPL\KTPPLW\CQLGQH
						SFMHSFLVIPHL/CPIFLIRGRNIL
İ						TKLS\ASLTIPGSTANISLVAFFP
		İ		ļ		NPKPPLCPLTSPQYHPLPQDLPS
						A
21046	51414	Α	21173	2	245	LTALLRKSLDPRLISTGENTQTA
						ETMGPLSAPPCTQHITW\KGLLL
1						TAETPKPSISSSNLNPRGGHGRL
		l				CALILLILEDSG
21047	51415	A	21174	3	324	
21048	51416	Α	21175	3	552	
21049	51417	A	21176	3	435	
21050	51418	Α	21177	3	1459	GAVRTWGRGFQTEKCQASLLN
						FWNPPTTAQVTIEAEPTKVSKG
				]		KDVLLLVHNLPQNLAGYIWYK
						GQMKDLYHYITSYVVDGQIIIY
			1			GPAYSGRETVYSNASLLIQNVT
						REDAGSYTLHIVKRGDGTRGET
		l				GHFTFTLY\LIPWTLLYL*RSLA
		l				QVSHYETPGKTAPPCGSTA**L
						N*PPDLTLFSPVLFLLKYPVPGQ
		l			·	AFQYPKGLKTIGSSITHL*DVLG
						KGSCRENIPRAAK*D*N*EDSST
						ACSK*GPQVETPKPSISSSNLYP
		l				REDMEAVSLTCDPETPDASYL
		l				WWMNGQSLPMTHSLQLSKNK
		ŀ				RTLFLFGVTKYTAGPYECEIRN
						PVSASRSDPVTLNLLPKLPKPYI
						TINNLNPRENKDVLAFTCEPKS
						ENYTYIWWLNGQSLPVSPRVK
			1			RPIENRILILTQCSREMKQGPY\
			1			QCEIQ\DRYGG\IRSYPVTLECP/
						HITTKHSGLYACSVRNSATGME
	1	L			<u></u>	SSKSMTVKVSAPSGTGHLPGLN

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21051	51419	Α	21178	3	1445	IRADPAHELENCSCPGKRLSTE
						RGRTAQLTAVLREFLDPRLISTE
ļ						ENTQAAETMGTLSAPPCTQRIK
						WKGLLLTASLLNFWNLPTTAQ
						VTIEAQPPKVSEGKDVLLLVHN
		l				LPQNLTGYIWYKGQMRDLYHY
						ITSYVVDGEIIIYGPAYSGRETA
				İ		YSNASLLIQNVTREDAGSYTLH
						IIKGDDGTRGVTGRFTFTLHLET
						PKPSISSSNLNPRETMEAVSLTC
						DPETPDASYLWWMNGQSLPMT
1						HSLKLSETNRTLFLLGVTKYTA
	İ					GPYECEIRNPVSASRSDPVTLNL
		=				LPKLPKPYITINNLNPRENKDVL
						NFTCEPKSENYTYIWWLNGQSL
						PVSPRVKRPIENRILILPSVTRNE
						TGPYQCEIRDRYGGVRSDPVTL
i						NVLYGPDLPRIYPSFTYYRSGE
						VLYLSCSADSNPPAQYSW\TINE
		1				KFQL\QGQKLFIP\HITTKHSGLY
						V\CSVRNSATGKESSKSMTVEV
						SGKWIPASLAIGF
21052	51420	Α	21179	3	1380	TAQLTAVLREFLDPRLISTEENT
						QAAETMGTLSAPPCTQRIKWK
						GLLLTASLLNFWNLPTTAQVTI
						EAEPTKVSEGKDVLLLVHNLPQ
						NLTGYIWYKGQMRDLYHYITS
						YVVDGEIIIYGPAYSGRETAYSN
						ASLLIQNVTREDAGSYTLHIIKG
		l				DDGTRGVTGRFTFTLHLETPKP
ļ		İ				SISSSNLNPRETMEAVSLTCDPE
Ē		ŧ				TPDASYLWWMNGQSLPMTHSL
į						KLSETNRTLFLLGVTKYTAGPY
						ECEIRNPVSASRSDPVTLNLLPK
						LPKPYITINNLNPRENKDVLNFT
						CEPKSENYTYIWWLNGQSLPVS
						PRVKRPIENRILILPSVTRNETGP
						YQCEIRDRYGGIRSDPVTLNVL
						YGPDLPRIYPSFTYYRSG\EVLL
						LVLFCGTLNPP\AQYSWTINEKF
						QLPGQKLFIRHITTKHSG\LYVC
		İ				SVRNSATGKESSKSMTVEVSAY
						SSSINYTSGNRN
	<u> </u>	<u> </u>	<u> </u>	<u> </u>	<u> </u>	

SEQ ID	-	1	SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21053	51421	Α	21180	2	1341	TEGGRTAQLTAMLRQLLDPRLI
						STEENTQAAETMGPLPAPSCTQ
						RITWKGLLLTASLLNFWNPPTT
1						AEVTIEAQPPKVSEGKDVLLLV
						HNLPQNLPGYFWYKGEMTDLY
						HYIISYIVDGKIIIYGPAYSGRET
						VYSNASLLIQNVTRKD\AGT\YT
						LHIIK\RGDETREEIRHFTFTLYL
						ETPKPYISSSNLNPREAMEAVR
						LICDPETLDASYLWWMNGQSL
						PVTHRLQLSKTNRTLYLFGVTK
						YIAGPYECEIRNPVSASRSDPVT
İ						LNLLPKLPIPYITINNLNPRENK
1						DVLAFTCEPKSENYTYIWWLN
						GQSLPVSPGVKRPIENRILILPSV
<u>l</u>						TRNETGP\YQCEIRDRYGGIRSY
						PVTLNVLYGPDLPRIYPSFTYYR
						SGENLDLSCFTESNPPAEYFWPI
						\*GKFQQSGQKLFIPQITRNHSG
					<u></u>	LYACSVHNSAAGKEISVSLFRH
21054	51422	Α	21181	1	1532	MTQFPMDKRQGRSRSVAITNT
1		1				AVTSLEWVPFHAFAIPRSTLVS
1	-	ŀ				QQRLSTERGRTAQLTAVLREFL
1						DPRLISTEENTQAAETMGTLSA
						PPCTQRIKWKGLLLTASLLNFW
						NLPTTAQVTIEAEPTKVSEGKD
		1				VLLLVHNLPQNLTGYIW\YKGA
		1				MR\DLYHYVTSYVVDGQIIKYG
						\PAYSGRETAYSNASLLIQNV\T
						RED\AGSYTLHIIKGDDGTRGVT
						GRFTFTLHLETPKPSISSSNLNPR
						ETMEAVSLTCDPETPDASYLW
						WMNGQSLPMTHSLKLSETNRT
						LFLLGVTKYTAGPYECEIRNPV
						SASRSDPVTLNLLPKLPKPYITI
						NNLNPRENKDVLNFTCEPKSEN
1						YTYIWWLNGQKPP/VSSPRVKR
ĺ						PH*KTGSLIPTPVFTRNETGP\YQ
						CEIRAR\YGG\IRSDPVTLNVLY
			}			GPDLP\RIYPSFTYYRSGGKSSY
			1			LSCFCGTSNPP\AQYSW\TIN*K
		1				VFSLPGTKALYSRHITTKHSGL
						YVCSVRNSATGKESSKSMTVE
						VSDWTVP
21055	51423	В	21182	28	29751	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
21056	51424	A	21183	158	1660	EASLLEDDDGMVYLRPLTSHQ
				1.00		YRTAQLRAMLRKFLDPRLSSTE
						ENTQAAETMGPLSAPPCTQRIT
						WKGLLLTASLLNFWNPPTTAQ
						VTIEAEPTKVSKGKDVLLLVHN
						LPQNLAGYIWYKGQMKDLYH
						YITSYVVDGQIIIYGPAYSGRET
						VYSNASLLIQNVTREDAGSYTL
						HIVKRGDGTRGETGHFTFTLYL
						ETPKPSISSSNLYPREDMEAVSL
						TCDPETPDASYLWWMNGQSLP
						MTHSLQLSKNKRTLFLFGVHK
						VHCRDPMECEIREPS*VASRS*P
						SSPWNLLPKL\PKPYITI\NN\LNP
						REKKDVLAFT\CDP\KSRNYTYI
						WWLNGQSLPVSPRVKRP\IENR
						DPSFLP\SVTRNETGP\YQCEIRD
		1				\RYGG\IRSYPVTLNVLYG\PDLP
						RIYPFISPIYHFREENLRTLSCFC
'						GTFNPPGRVFLGTINGGSFQVIQ
						GQKALYSPRITTKHSGA/LYAW
						LCSVTSRPLAWESSQIPWTV\KV
!						SGSFQGTGTSSWGLNPL
21057	51425	Α	21184	927	1378	QPPTLRSWDCWFAWGKRQLM
						SSP/STEPQVCFTIEGHEIDFLLG
						TGMAFSVLISCPG*LSSRSVTIR
						GILG*PVTRYFSHLLSCNWETLL
						FSHNAKTQGGGTLTLPKPSKSH
		1				RPWAVRGLAPGPAAAVLNFSP
						GLSCLPTGQGLGPAAHHA
21058	51426	Α	21185	1	221	ILLAQLESQDGDILRQRDEFDLL
						VAGEICRFSALIEELE\EKNERPA
						RELLTVRPEPEPCPTCAVLRLLH
		1				LCLA

SEQ ID	1 -	1	SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				Sequence		
21059	51427	Α	21186	335	1963	VCDPALPPWAKQREWALEGSH
						SSRTSMGLLVASVTSLGRWKF
						NCPI/CVQGTLKEAGSLSNCG/H
						KNFCRACLT\RYCEIP\GPD\LEE
1						SP\TCP\LCKEPFRP\GSFRPNWQ
						LANVVENIERLQLVSTLGLGEE
						DVCQEHGEKIYFFCEDDEMQL
						CVVCREAGEHATHTMRFLEDA
				•		A\APYREQIHKCLKCLRKEREEI
						QEIQSRENKRMQVLLTQVSTKR
						QQVISEFAHLRKFLEEQQSILLA
				1		QLESQDGDILRQRDEFDLLVAG
						EICRFSALIEELEEKNERPAREL
	1					LTDIRSTLIRCETRKCRKPVAVS
						PELGQRIRDFPQQALPLQREMK
						MFLEKLCFELDYEPAHISLDPQ
						T\SHPKLPLVPRTIQR\AQFS*QM
						GRTSPDNPQ\RF\DRATCVSRPT
						LGITGGRHTWVVSIDLAHGGSC
						TVGVVSEDVQRKGELRLRPEE
	1					GVWAVRLAWGFVSALGSFP\TR
						LTLKEQPRQVRVSLDYEVGWV
						TFTNAVTREPIYTFTASFTRKVI
						PFFGLWG\RGS\SFSLSSLRRSSY
						LLSSKYRTHINPSTMWT
21060	51428	A	21187	217	397	
21061	51429	Α	21188	529	1526	YCGRVGELAAGELVFLWSRVG
	1					PQNPRSSRLPPPTLAMFLTRSEY
						DRGVNTFS\PEGRLFQVEYDIEA
	1					IKLGSHSQLAPRHREGV\ALAV\
		ł				EKRIYLPHLMGAQAGI*GKLLG
		1				DLMLHIGLLPIEWGLICLMLKT
		l				FNSIKSRVGRHR/YPWGSPYNG
		1				GH*QWESFEPKLCPNLGFSSFG
						EEDAESRVPMS\RPLGSSHLFIW
						EELYEKGTPSLFP*GTPFWGPFV
						PVWMARSQFGSGFQRGAPRAP
						LPRKFPPTSPMTFERSPSSPSPHP
		ŀ				SLKQVMGGRKLNANQTIEPRPT
						VPALAENFPHCSQRKKLEEVIQ
						GHFKESWILRTSLGTISVLNNVP
21062	51430	Α	21189	211	382	TSVSPSVPPLPC*RHPGEPQTGN
					[	NASNGTSSSRDESRPPVRVVPPP
						QTAVAITLSTA
21063	51431	A	21190	3	96	FFFPPQKYD*DR*EHPRFHKTLR
		1				NWRVGAQ
21064	51432	Α	21191	1	1320	
21065	51433	A	21192	1	339	
	1			<del></del>	<del></del>	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21066	51434	Α	21193	289	3694	MAFVPVIPESYSHVLAEFESLDP
		l				LLSALRLDSSRLKCTSIAVSRK
						WLALGSSGGGLHLIQKEGWKH
						RLFLSHREGAISQVACCLHDDD
1						YVAVATSQGLVVVWELNQERR
						GKPEQMYVSSEHKGRRVTALC
						WDTAILRVFVGDHAGKVSAIK
		ŀ				LNTSKQAKAAAAFVMFPVQTIT
		ŀ				TVDSCVVQLDYLDGRLLISSLT
						RSFLCDTEREKFWKIGNKERDG
		1				EYGACFFPGRCSGGQQPLIYCA
					•	RPGSRMWEVNFDGEVI
21067	51435	Α	21194	2	128	
21068	51436	Α	21195	1041	2238	LLHPAQRSLSKDGTKESLEDAA
	]					LMGEEGKPEINQQLSLESMELD
	•					ELALEKYPIAAPLVPYPEKSSED
	ŀ	ŀ				GVGNPEAKILSGTPTYKRRVISL
						LVTIENHTPLVELSEYLGTNTLS
						EILDSPWEGAKNVYKCPECDQ
						NFSDHSYLVLHQKIHSGEKKHK
						CGDCGKIFNHRANLRTHRRIHT
						GEKPYKCAKCSASFRQHSHLSR
						HMNSHVKEKPYTCSICGRGFM
						WLPGLAQHQKSHSAENTYEST
						NCDKHFNEKPNLALPEETFVSG
						PQYQHTKCMKSFRQSLYPALSE
	1					KSHDEDSERCSG\CGDNFFSFSK
						FKPLQCPDCDMTFPCFSELISHQ
						NIHTEERPH*\CKTCEESFALDSE
		1				LACHQKSHMLAEPF\KCTVCGK
						TFKSNLHLITHKRTHIKKHHVK
21069	51437	C	21196	156	389	

SEQ ID NO: of peptide sequence sequence sequence sequence sequence 21070 51438 A 21197 1 2466	of last Amino acid sequence (X=Unknown, > acid *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
sequence	deletion, \=possible nucleotide insertion)
21070 51438 A 21197 1 2466	
210/0  31438  A  21197  1  2400	MI ODI CDAMICI CUA CHDWCVI
	MLSRLFRMHGLFVASHPWEVI
	VGTVTLTICMMSMNMFTGIAG
	LFTIFSSFVFSTVVIHFLDKELTG
	LNEALPFFLLLIDLSRASTLAKF
	ALSSNSQDEVRENIARGMAILG
	PTFTLDALVECLVIGVGTMSGV
	RQLEIMCCFGCMSVLANYFVF
	MTFFPACVSLVLELSRESREGR
	PIWQLSHFARVLEEEENKPNPV
	TQRVKMIMSLGLVLVHAHSRW
	IADPSPQNSTADTSKVSLGLDE
	NVSKRIEPSVSLWQFYLSKMIS
	MDIEQVITLSLALLLAVKYIFFE
	QTETESTLSLKNPITSPVVTQKK
	VPDNCCRREPMLVRNNQKCDS
	VEEETGINRERKVEVIKPLVAET
	DTPNRATFVVGNSSLLDTSSVL
	VTQEPEIELPREPRPNEECLQIL
	GNAEKGAKFLSDAEIIQLVNAK
	HIPAYKLETLMETHERGVSIRR
	QLLSKKLSEPSS\LQYLPYRGY\
	NYSLW*WGACCENVIGYMPIP
	VGVAGPLCLDEKEFQVPMATT
	EGCLVASTNRGCRAIGLGGGAS
	SRVLADGMTRGPVVRLPRACD
	SAEVKAWLETSEGFAVIKEAFD
	STSRFARLQKLHTSIAGRNLYIR
	FQSRSGDAMGMNMISKGTEKA
	LSKLHEYFP*NCRF*AV\SGNYC
	TDKKPAAINWIEGRGKSVVCET
	VIPAKVVREVLKTTTEAMIEVN
	INKNLVGSAMAGSIGGYNAHA
21071 51439 A 21198 236 3022	GEGSFRSATALTGARLSVGAGV
	RWPLVRSGGSKDSVATMLSRL
	F\RMHGLEVASHPWEVILGTVT
	LTICMMSMNMFTGNNKICGWN
	YECPKFEEDVLSSDIIILTITRCIA
	ILYIYFQFQNLRQLGSKYILGIA
	GLFTIFSSFVFSTVVIHFLDKELT
	GLNEALPFFLLLIDLS\RASTLA
	KFALSSNSQDEVRENIARGMAI
	LGPTFTLDALVECLVIGVGTMS
	GVRQLEIMCCFGCMSVLANYF
01070 51440	VFMTFFPAC
21072 51440 A 21199 1 450	CKSRGSNLRVHFKNTRETAQAI
	KGMHIRTATKYLKDVTLQKQC
	VPFRRYNGGVGRCAQAKQWG
	WTQGRWPKKSAEF\LVIEHIQV
	NKAPKMRRRTYRAHGRINPYM
	SSPCHIEMILTEKEQIVPKPEEEV
	AQKKKISQKKLKKQKLMARE

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21073	51441	Α	21200	1	876	
21074	51442	Α	21201	1	1725	
21075	51443	A	21202	278	889	SVKMVRYSLDPGGTPRKSCK/S QRGSNLRVPFKDHS*KLPQAHQ RVCHIRKSPTKY\LKDVHLTRN QCVPIPDRYNG*QLGQVCRRHK QMGPGTTKGR\WPQKGVLKFL PAHALKTAE\SNAEL\KGLDVDS LVIEHI\QVNKAP\KM\RRRTYR AHGRINPYMSSPCH\IEM\ILTEK EQIVPKPEEEVAQKKKISPEETE RNKNLLARE
21076	51444	Α	21203	1	1923	
21077	51445	Α	21204	1	280	
21078	51446	Α	21205	1	109	
21079	51447	Α	21206	1	638	
21080	51448	Α	21207	1	396	
21081	51449	A	21208	26	465	HGVGCRDGLPMECRFLCQIFM AFMQMNSYCARHKKATRRTV* /CRNQDMTESSQKWTGKGSAV EHLVESGEWTMHPARPVSACR CPSDCHVPLSPGACWPKLECLS EGSSYISDGKEYLFYLNVCGET EIQFCNKKQAAVCQVKKSDTS QVKAAGRYHNQTLRPSESSWS LLFVVLVFFPGVHSPFSLDEALT LCPQNCLPPGTVSAPLNCVYPL ALSAAVALAAGSELMECHRLV NSLEVSTAAQGQSSKTQFYACL CRLSTHLWKYSEESGRRLVCSF YQPSELNPALNLSFLVGTRLQE GAALLFIQSGFTIV
		A		1		
21083	51451	A	21210	217	453	
21084 21085	51452 51453	A A	21211 21212	1	840 663	-
21086	51454	A	21213	176	957	DEPIGVPLLSAPPAGTTKIKADQ LHLQGAAPKICPCLQDGSLGQH YCPFVSFSRYYINLCQKIYKGPL GCSERASICRRTTTGDVQVLG\L VHTQK\LGVIGPRVVVTTSKRY PCGGNKTASSVIELTCTKPVGR PAFKRFEIDSCTYYFSWDSGAA CAVKPQEVQMVNGTITNPINGK SFSLGDIYFKLFRASGDMRTNG DNYLYEIQLSSITSSRNPACSGA NICQVKPNDQHFSRKVGTSDKT KYYLQGNPWLPTKFHI

SEQ ID NO:	SEQ ID NO: of peptide sequence	i	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21087	51455	A	21214	3	7616	SPVEPRSPRAPAPSPPPPLPWRR DPVPGAAPSSRAPLASRPPRSSG PGAMGAAAGRSPHLGPAPARR PQRSLLLLQLLLLVAAPGSTQA QAAPFPELCSYTWEAVDTKNN VLYKINICGSVDIVQCGPSSAVC MHDLKTRTYHSVGDSVLRSAT RSLLEFNTTVSCDQQGTNHRVQ SSIAFLCGKTLGTPEFVTATECV HYFEWRTTAACKKDIFKANKE VPCYVFDEELRKHDLNPLIKLS GAYLVDDSDPDTS
21088	51456	A	21215	2	184	VNCQPLGMISLMKR\PPGFSPFR SSRIGEIKEETTSHLRSCEYKGR PPKAGAEPASEREVS
21089	51457	Α	21216	1	203	
21090	51458	Α	21217	l	1107	
21091	51459	Α	21218	l I	915	
21092	51460	Α	21219	l	1932	
21093	51461	Α	21220	2	746	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21094	51462	A	21221	1	2157	MGFPLRPSPAHLSSGRQQFPAK
21094	31402	^	21221	L	2137	EERLLISLTQESQSEEIDCNDKD
	1					LFKAVDAALKKYNSQNQSNNQ
						FVLYRITEATKTVGSDTFYSFK
						1
						YEIKEGDCPVQSGKTWQDCEY
						KDAAKAATGECTATVGKRSST
						KFSVATQTCQITPGGWFILWHC
			Ė			PGGKLTISDYPNSSRVLHCTQC
						VQGLNRRTRRHKTCPPLQAAS
						QIEERIGFEGRPGLESFRAPCLL
						ASPFPCLEKGAEGPVVTAQYDC
						LGCVHPISTQSPDLEPILRHGIQ
						YFNNNTQHSSLFMLNEVKRAQ
						RQAAP/GREFSSMDTFEGYEGA
						EDMEKLAGPFPVG*QSRNLFLQ
						IRFPGEERKHPGRKRRTR*GSN
						GDRQHQEGACWGVHAG*LLR
						GRRLVSRRGRGARGPPPRRAAL
		1				PAVAPAPRLQSARIPERGDGDD
						GVRRWGHREAIGS*GQS/LS/CN
						GIIILIKEERLPCPFHCISLMKRPP
						GFSPFRSSRIGEIKEETTVSPPHT
						SMAPAQDEERDSGKEQGHTRR
						HDWGHEKQRKHNLGHGHKHE
						RDQGHGHQRGHGLGHGHEQQ
ľ		i				HGLGHGHKFKLDDDLEHQGGH
						VLDHGHKHKHGHGHGKHKNK
						GKKNGKHNGWKTEHLASSSED
	İ					STTPSAQTQEKTEGPTPIPSLAK
						PGVTVTFSDFQDSDLIATMMPPI
		1				SPAPIQSDDDWIPDIQIDPNGLSF
						NPISDFPDTTSPKCPGRPWKSVS
21095	51463	A	21222	207	424	HHINCVLLYLQSHLRS\CEYQG\
21093	21403	\ <u>^</u>	21222	207	747	RPPKAGAEPASE\REVSLTNWA
						l :
						ESSTPRHIAPTTSASNLEMEGQE
			<u> </u>	<u> </u>		ERWDRI

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21096	51464	A	21223		1487	IMKLITILFLÖSRLLLSLTQESQS EEIDCNDKDLFKAVDAALKKY NSQNQSNNQFVLYRITEATKTV GSDTFYSFKYEIKEGDCPVQSG KTWQDCEYKDAAKAATGECT ATVGKRSST\KFSVATQT\CQITP AEGPVV\TAQLRTCLG\CVWHPI IQRQSPDLEPILRHGIQYFNNNT QHSSLFTLNEVKRAQRQVVAG LNFRITYSIVQTNCSKENFLFLT PDCKSLWNGDTGECTDNAYIDI QLRIASFSQNWD\IYPGKDVVQ PPTKICVGCPRDIPTNSPELEETL THTITKLNAENNAISYFKIDNVK KARVRVVAGKKYFIDFVAR\ET TCSK\ESNEELTESCETKKLG\Q RLG\CNAE\VYVVPW\EKKIYPT\/ VFNCSTTGEWIFT*WKRPPRVF SPFRS\SRIGEIKEET\TSHLRSCE YKGRPPK\AGAEPASEREVS
21097	51465	С	21224	21	281	
21098	51466	A	21225	204	441	QISQVLIICNQK*VILTGNFLDT YF*TTFINKKIDELRLCA/CAVA HSYNTTLCKKPRWKDCLRPAV QDQPVQHSETLSL
21099	51467	A	21226	333	604	SSDRITKPPSAEFLPLQRTFLHSL QDTVIELLH*DLETPPLTVLQNH PSLPRPIPAPPTLCPQTRWPHLR GHQGILAASHILETALAAIF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21100	51468	A	21227	846	3269	GFMEVLRDPIKKNSSESKPAQS
	ŀ					GFSRGNSPLSCPESVEASPAVNE
						KSVYSTHNYGTTQRHGCRGLP
						YADHNYGAPPPPTPPASPPVQTI
						IPRSDLNGLPSPVEERCGDSPNS
						EGETVPTWCPCGLSQDGFLLNC
						DKCRGMSRGKVIRLHRRKQDN
						ISGGDSSATESWDEELSPSTVLY
						TATQHTPTSITLTVRRTKPKKR
Ī						KKSPEKGRAAPKTKKIKAFREG
						SRKSLRMKNSPSEAQNLDENTT
						EGWENRIRLWTDQYEEAFTNQ
			1			YSADVQNALEQHLHSSKEFVG
ļ						KPTILDTINKTELACNNTVIGSQ
						MQLQLGRVTRVQKHRKILRAA
						RDLALDTLIIEYRGKVMLRQQF
						EVNGHFFKK\PYP\FVLFLLQNL
						NGVEMCVDARTFGNDARFIRR
						SCTPNAEVRHMIADGMIHLCIY
						AVSAITKDAEVTIAFDYEYSNC
						NYKVDCACHKGNRNCPIQKRN
						S*CYRTATPTTSSKPTHHWSRD
		1				*T*KS\RRKELEMEQQNEASEEN
						NDQQSQEVPEKVTVSSDHEEV
						DNPEEKPEEEKEEVIDDQENLA
						HSRRTREDRKVEAIMHAFENLE
		ŀ		:		KRKKRRDQPLEQSNSDVEITTT
						TSETPVGEETKTEAPESEVSNSV
						SNVTIPSTPQSVGVNTRRSSQA
						GDIAAEKLVPKPPPAKPSRPRPK
						SRISRYRTSSAQRLKRQKQANA
						QQAELSQAALEEGGSNSLVTPT
21101	51469	Α	21228	2	197	CRPPSPRPVRAARPARPQASFSR
						EPPPRRMAP*SKRRRQSARGPV
						RPSRVWARIKAKPYSCCCQL
21102	51470	Α	21229	4	488	GSYDINRQDTFQKDRQRVGPSG
						ASQKTEGFLLRLLYSGLIIAHCS
						LEILGRNDPPASASKVAETTDS
						GSLSLANT*PPTPR/VTSQTANA
						TSQAAPALPLGSVPGRRAADPR
						PLLAWAPAPPGPGTSRNTHPAP
						DGLDMLVPIGDGVLGGNCFRN
						QCKSVSTS
21103	51471	Α	21230	124	293	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	I .	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
21104	51472	Α	21231	239	1393	IYPQISFQKKFVMFFFSEALCTA
ŀ						RAEECDPIGLLNNPRTSRYSRL
						KPNSSLGGSESRKTQMGPLPPP
						PKSPDGARPVPGCIKEAGPARG
						RSRLAPNPGALRRG/VGLGR*S
						ARI*GLVSKAAAGQGEARRGE
						AT*ARGALGRTTAGSFNAGWL
						TLDGARQQPGRQTPTRIPSCVA
						DNAAGPSRTTTPD/EALRPLRG
		i				GPACTAPASFDSRAPYSASSAPS
						EGRWPSRGFPPIS/PLYSQLPETR
						SHEWQAASKIQPRKASTSQ*SA
						RMEAT*QQWAGPPAEP*SLWFP
						RLARRSVVHWRTPY/ISSSITSLP
						PVAAGIPEDSGPGSTQSSSRPGN
		1				TSSSSRPPPPPPPPPPP/QPLPPPPA
						GLASPRPVRARRPARPQASFSR
						EPPPRRMAP
21105	51473	Α	21232	107	234	
21106	51474	Α	21233	224	227	LPQPGPGRRRACLRQASPRQYS
						RLPGRPPPSG*SW
21107	51475	Α	21234	67	607	QRSWAGPGAGPEAGTRPPARG
						RRRQPGNVDPRRRAPQLRSQM
	İ					QVAMARATTATGNRLWPGLLI
		1				MLGSVCHRGSPCGLSTHVEI\LL
		1				LEHQDAYQAGIVFPDCFYPSIC
						KGGKFHDVSESTHWTPFLNAS
•						VHYIRENYPLPWEKDTEKLVAF
1						LFGITSHMAADVSWHSLGLEQ
						GFLRTMGAI
21108	51476	A	21235	54	193	
21109	51477	Α	21236	1275	1764	STKCQYFSHLLLLFSASESKQLL
						LYSAFFRYVPVKDLLGIYEKLYI
		1				CFISLLQTGL*RYQE*YHIHFIFS
						FSLP*LYPTYSTKSPFLVEQFQE
						YFLGGLDDR*NSCL*RSR*MAN
		1				SFDCFLSDSVLQGVVFSVVTSL
						FLFLFNSSKMQKNDFHRNLTTS
21110	51478	Α	21237	297	428	QCQRGPGCSRGHGI*I*EHWTA
		L				FGLLDLRQWFARGSWASGHGP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
21111	51479	A	21238	3247	5958	  QRSWAGPGAGPEAGTRPPARG
						RRROPGNVDPRRRAPOLRSOM
						QVAMARATTATGNRLWPGLL
						MIVMASLCHRGSSCGLSTHIEIG
						HRALEFLHLHNGHVNYKELLL
				ŀ		EHQDAYQAGTVFPDCFYPSLC
						KGGKFHDVSESTHWTPFLNAS
						VHYIRENYPLPWEKDTEKLVAF
						LFGITSHMVADVSWHSLGIEQG
						FLRTMGAIDFHGSYSEAHSAGD
						FGGDVLSQFEFNFNYLARRWY
						VPVKDLLGIYEKLYGREVITEN
						VIVDCSHIQFLEMYGEMLAVSK
					:	LYPSYSTKSPFLVEQFQEYFLG
						GLDDMAFWSTNIYHLTSFMLE
						NGTSDCSLPENPLFIACGGQQN
						HTQGSKMQKNDFHRNLTSSLT
						ENIDRNINYTERGVFFSVNSWT
						PDSMSFIYKALERNVRTMFIGG
						SQLSQKHVSMPLRHHYFLSFPY
						ARLGW\AMTSADL\NQDG\HGD
						LRGWGAPGLTAAPAHIPHSGR
						VYLIYGNDLGLPPVDLDLDKEA
						HRILEGFQPSGRFGSALAVLDF
						N\LDGV\PDLAVGAPSVGSEQLT
	Ì					YKGAVYVYFGSKQGGMSSSPN
						ITISCQDIYCNLGWTLLAADVN
		l				GDSEPDLVIGSPFAPGGGKQKG
						IVAAFYSGPSLSDKEKLNVEAA
						NWTVRGEEDFSWFGYSLHGVT
						VDNRTLLLVGSPTWKNASRLG
						HLLHIRDEKKSLGRVYGYFPPN
21112	51480	Α	21239	2	403	VVAEEDTELRDLLVQTLENSGV
						LNRIKAELRAGVFLALEEQEKV
	1					ENKTPLVNESLKKFLNTKD\LQ
	-					GLEGRENLARDLGIIEAEGTVG
		1				GPLLLEVIRRCQQKEKGPTTGE
		1				GALDLSDVHSPPKSPEGKTSAQ
21113	51481	Α	21240	3	1699	
	<u> </u>			J	I	<u> </u>

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21114	51482	A	21241	3	1812	APRLKMGLHPNKPIVSQSRVVN DGQCKLHAAVDSRDPPDGTKR RVSSGGTGLPGPGRGPGLPAGA WLSAEGRGAGGSGDCGPSSGS AEPTQRPPRPSALPGAIPPLHLG LAPFRLASEAVAEQAPPPAPTLS FGAQWSGAPQGSVSSVNPEGA GPARCASGRGFGCVGRVARAR SVPCCGRWPLARLRRLSWRSK MAATAAAVVAQEDTELRDLL VQTLENSGVLNR\IKAELRAAV FLALEEQEKVENKTPLVNESLK KFLNTKDGRLVASLVAEFLQFF NLDFTLA\VFQPETS/TIF*LLF*L QGLEGRENLARDLGIIEAEGTV GGPLL\LEVIRRCQQKEKGPTTG EGALDLSDVHSPPKSPEGKTSA QTTPSKIPRYKGQGKKKTSGQK AGDKKANDEANQSDTSVSLSE
						PKSKSSLHLLSHETKIGSFLSNR TLDGKDKAGLCPDEDDMEGDS FFDDPIPKPEKTYGLRKEPRKQ AGSLASLSDAPPLKSGLSSLAG APSLKDSESKRGNTVLKDLKLI SDKIGSLGLGTGEDDDYVDDFN STSHRSEKSEISIGEEIEEDLSVEI DDINTSDKLDDLTQDLTVSQLS DVADYLEDVA
21115	51483	A	21242	419	651	TNIFSRLNPQTKNVFCARQLSS* SVTI*GVLGQAVTRYFSQALSC DWGTLLFSRAFLIMPESPTPLLG RDILAKAGA
21116	51484	A	21243	555	888	RTSSRRTPNPCSRIRKNYFKFH MEPKKSP/S*PRQC*AKRTKLEA SCYLTSNYTTRLQ*PKQHGWG TIRSFSSHEAIFQTITWGKTLDN TRLSRTEVPAAFHSALCPWFIET
21117	51485	Α	21244	728	1428	
21118	51486	A	21245	238	505	INVGRHHVILQGLSCPGQLSSRS VTI*G\VLGQAVTIYFSQPLSCD WRTLLFSHVFLIMPESPTPLLGR DILLYVCIGRFPIFRADCRT
21119	51487	A	21246	940	1434	QLAPCTLTEPRVCLTIEGQEVN CFLDTRAAF\VLLSCPRQLSSRS VAIRGVLGQAVT*YFSQPLSCD WGTLRFSHAFLIMPKSPTPLLG RDILAKAGAIIHEKRKAFMRKR EIKKQARSSKVHTSAASSCART APLGESFAEHLIKIADFTPPSSSS PCTTLFFS

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21120	51488	В	21247	747	1138	
21121	51489	A	21248	1685	2051	SSSRTEGARGKRQPMPSP/STEP RVCLTIEGQEVNCLLDTGAAFS VLLSCPGQLSSRSVTI*GVLGQP VTRYFPQPLSCDWGTLLFSNAF LIMPESPTPLLGRGILAKAGAIIH
21122	51490	A	21249	2	1014	HAVLVTAAL  GGAGPVVLRRRAACSPGPAPR  AARPPAMEPGPDGPAASGPAAI REGWFRETCSLWPGQALSLQV EQLLHHRRSRYQDILVFRSKTY GNVLVLDGVIQCTERDEFSYQE MIANLPLCSHPNPRKVLIIGGGE GG\VL\REVVKHPSVESVVQCEI DEDVIQVSKKFLPG\MAIGYSSF EADTYMWGDGF*VSWKQKQ\D AFDVI\IIDSSDPMGPA\ESLFKES YYQLMKTA/LLREDGVL\CCQG \ECQWLH\LDLIKE\MRQFC\QSL FPV/VLAYAYCTIPT/YTPSGQIG FMLCSKNPSTNF\QEPVQAVDN SSKVAKMQLK\Y\YNSERATAA AFVLPE\FARQGLE
21123	51491	A	21250	3	418	FSYEKLLNWQEFAVDLEHHSY RSFLGLTCLKQISTRTEDFIIDTL ELRRDMYILNESLTDPAIVKP\L AEEMLSYARDDTHYLLYIYDK MKLQMRERGNRQTVQLHVVW QRSRDICLKKFIKPI\FTDESYLE LYRKLKK
21124	51492	A	21251	1	1704	
21125	51493	Α	21252		3104	MFDTHQAARLLNLGRHSLDHL LKLYCNVDSNKQYQLADWRIR PLPEEMLSYARDDTHYLLYIYD KMRLEMWERGNGQPVQLQVV WQRSRDICLKKFIKPIFTDESYL ELYRKQKKHLNTQQLTAFQLL FAWRDKTARREDESYGYVLPN HMMLKIAEELPKEPQGIIACCN PVPPLVRQQINEMHLLIQQARE MPLLKSEVAAGVKKSGPLPSAE \RENVKSQPVQVHL*QRPVLTA GRKLLPACTFYMPACWGL

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21126	51494	A	21253	2	2775	QALARRA VQAEKMAPPSTREP RVLSATSATKSDGEMVLPGFPD ADSFVKFALGSVVAVTKASGG LPQFGDEYDFY\RSFPG\FQAFC ETQGDR/LCFQCMSRVMQVPM GC\RSN\IKDRSKV\TELEDKF\D LL\VDAN\DVILERVGILLDEAS GVNKNQQPVLPAGLQVPKTVV \SSWNRKAAEYGKKAKSETFRL LHAKNIIRPQLKFREKIDNSNTP FLPKIFHSNPMVQ\KPLPQALSK
21127	51495	A	21254	3	2668	ERRERPQDRPE  QALARRAVQAEKMAPPSTREP  RVLSATSATKSDGEMVLPGFPD  ADSFVKFALGSVVAVTKASGG  LPQFGDEYDFYRSFPGFQAFCE  TQGDRLLQCMSRVMQYHGCRS
						NIKDRSKVTELEDKFDLLVDAN DVILERVGILLDEASGVNKNQQ PVLPAGLQVPKTVVSSWNRKA AEYGKKAKSETFRLLHAKNIIR PQLKFREKIDNSNTPFLPKIFIKP NAQKPLPQALSKERRERPQDRP EDLDVPPALADFIHQQRTQQVE
						QDMFAHPYQYELNHFTPADAV LQKPQPQLYRPIEETPCHFISSL DELVELNEKLLNCQEFAVDLEH HSYRSFLGLTCLMQISTRTEDFII DTLELRSDMYILNESLTDPAIVK
:						VFHGADSDIEWLQKDFGLYVV NMFDTHQAARLLNLGRHSLDH LLKLYCNVDSNKQYQLADWRI RPLPEEMLSYARDDTHYLLYIY DKMRLEMWERGNGQPVQLQV
						VWQRSRDICLKKFIKPIFTDESY LELYRKQKKHLNTQQLTAFQL LFAWRDKTARREDESYGYVLP NHMMLKIAEELPKEPQGIIACC NPVPPLVRQQINEMHLLIQQAR
						EMPLLKSEVAAGVKKSGPLPSA ERLENVLFGPHDCSHAPPDGYP IIPTSGSVPVQKQASLFPDEKED NLLGTTCLIATAVITLFNEPSAE DSKKGPLTVAQKKAQNIMESFE
21128	51496	A	21255	98	286	HARPRHGPRLRCEVGLRCAPG YAGASARPTCECGDTRYCPQDP NQT*SRPVPIIAHRYCLTG
21129	51497	Α	21256	1020	1280	
21130	51498	В	21257	153	3804	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21131	51499	Α	21258	159	579	PNVRWCPS\CYRNQDDHELQIT
						HGNKILCGIVWKKIAQVEQMR
						KFSRLEKQRTHVLSCSDLQAVR
						RFRGVVSEGSFEMGEKVYKRK
						DMFQLSLLTEQHGDSTLGSGSS
						YTQMPAENGSLVHTLVTAAPPF
						FAEKPHSITAG
21132	51500	Α	21259	1	1290	MGASRSQKPGPSRDYSYTAGC
						DVDQRRRGAQKLSRSDTKRIV
						MSYGKKALKNDKFEIRAFSSSL
						EVRYRNQDDHELQIHTGTRFY
						VELCGFAVTRLRDPPMLRHAA
						MALLPWRPPDYSASLLPDWLP
						VPSSKVVITYPGFTSIWAFSNTE
						T/LCSEYKYYCEECRSKQE/AHK
						RRKV*KLPMILAP/HLKRFKYK
						DQLHRFTKLFTRASLRGPRDER
						LKRVSRQLFSLVVVTSRVFDVL
						LRILHRLGQYVSSEYYVALMCS
						EGHGPGAEDVTELLQSHEKTE
						WMRSCFFWMTKSGVFDGIYLW
						NAFNQEGEDLYKENCKTLMKE
						VEDAQINGKTSYAHGSEEYCK
						MSILPRILLQKISRDDCFKIVLKF
						CVDHQKTKKCCMITLKSSVVQ
						LAFTIKISVSEMCAREDFVVVIS
						FQHSEVPSRQLARCTFKEGYT
21133	51501	A	21260	3	142	
21134	51502	Α	21261	1	1368	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21135	51503	A	21262		2420	  MEEVAEAPQPIDPKPLATQPQSI
						IITTITTIITITTTTTTTTTTTTTTTTTTTTTTTTTT
						IITTTITITTTTTTTTTITITTTTTTT
						TITITTIITIITTTITTTITTTTTT
						VITIITTITTIITITTITPSSSSSPSS
						QADECWGEAEPLMSRQAHVW
						LRYRTGCREERPDQELTGSWG
						HGPRSTLVRAKAMAPPPPPLAA
						STPLLHGEFGSYPARGPRFALTL
						TSQALHIQRLRPKPEARPRGGL
						VPLAEVSGCCTLRSRSPSDSAA
						YFCIYTYPRGRRGARRKATRTF
						RADGAATYEENRAEAQRWAT
						ALTCLLRGLPLPGDGEITPDLLP
						RPPRLLLLVNPFGGRGLAWQW
						CKNHVLPMISEAGLSFNLIQTER
						QNHARELVQGLSLSEWDGIVT
						VSGDGLLHEVLNGLLDRPDWE
		į				EAVKMPVGILPCGSGNALAGA
						VNQHGGFEPALGLDLLLNCSLL
						LCRGGGHPLDLLSVTLASGSRC
						FSFLSVAWGFVSDVDIQSERFR
						ALGSARFTLGTVLGLATLHTYR
						GRLSYLPATVEPASPTPAHSLPR
						AKSELTLTPDPAPPMAHSPLHR
						SVSDLPLPLPQPALASPGSPEPL
						PILSLNGGGPELAGDWGGAGD
						APLSPDSTAGFTSWLSQGSSTLT
İ						RLRRGPRNSPILWAPTSHP*CPG
						RGLH\RGPPDHLLPPLGTPLPPD
						WVTLEGDFVLMLAISPSHLSAD
						LVAAPHARFDDGLVHLCCVRS
21136	51504	A	21263	573	1025	RKQSHEDEWRSTNKTVGHPSA
						RLYEPPWWAHTWTPPGGSKAP
						SDP/PLSHQLSPPAGQPHFASPEP
						TSKTPDPWPGNVFSSFPLPQSLP
						ELRHTLYLPMGTKKPPGSSMIF
						YT*AQYPGTNSSATICRLLSYKS
		i i				GQVYPNTGQAPPQQRRGR
21137	51505	A	21264	48	327	EELEALRR\QRLAELQAKHGDP
1213/	31303	'`	21207	10	521	GDA\AQQEAKHREAEMRNSILA
						QVLDQSARAR\VSEQGLIEILKK
-						VSQQTEKTTTVKFNRRKVMDS
						DEDDDY
21138	51506		21265	1	400	DEDDDI
		A	21265	786	905	VSEQGLIEILKK*CQLTEKTSTV
21139	51507	Α	21200	/80	000	KFNRRKVMDSDEDDDY
L	<u> </u>		<u> </u>	<u> </u>	<u>L</u>	MI NKKW A MINODEDDD I

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21140	51508	A	21267	907	1327	PRLLQRSTPSPWRTEELEALRR\ QRLADAAGQNTGDPGDAA\QQ EAKAPGQAEMRNSILAQVPGSS PAPAPRLSNLSHLVKP*KNLKP V\ENYPYTRLARYWTT*VMTGI RNQGFN*KSLKKVKAQ\QTEKT TTVKFNQKKK
21141	51509	В	21268	1	828	
21142	51510	Α	21269	317	2390	KWPKMNPQQQRMAAIGTDKE LSDLLDFSAMFSPPVNSGKTRP TTLGSSQFSGSGIDERG\GTTSW GTSGQPSP\SYDSSRGFTDSPHY SDHLNDSRLGAHEGLSPTPFMN SNLMGKTSERGSFSLYSRDTGL PGCQSSLLRQDLGLGSPAQLSS SGKPGTAYYSFSATSSRRPLH DSAALDPLQAKKVRKVPPGLPS SVYAPSPNSDDFNRESPSYPSPK PPTSMFASTFFMQDGTHNSSDL WSSSNGMSQPGFGGILGTSTSH MSQ\SSSYGNLHSHDRLSYPPHS VSPT\DINTSLPPMSSFHRGSTSS SPYVAASHT\PPINGSDSILGTK GNAAG\SSQTGDALGKALASIY SPDHTSSSFPSNPSTPVGSPSPLT GTSQWPRPGGQAPSSPSYENSL HSLQSRMEDRLDRLDDAIHVLR NHAVGPSTSLPAGHSDIHSLLG PSHNAPIGSLNSNYGGSSLVASS RSASMVGTHR\EDSVSLNG\NH SVLSSTVTTSSTDLNHKTQENY *RWPCKSQSGT\VVPQEIKT\EN KEKDENLHEPPSSDDMKSDD\* SSQKDIKVSSRGRTSSTNEDEDL NPEQKIEREKERRMANNARERL RVRDINEAFKELGRMCQLHLKS EKPQTKLLILHQAVAVIL\SLEQ QVRERNLNPKAACLKRREEEK VSAVSAEPPTTLPGTHPGLSETT
21143	51511	A	21270	2	153	QNASSPSKGSQLLTSKGTKLDG E*V**INRSRLQKVANNKLLWR KLKGQF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	1		Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence	ŀ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21144	51512	A	21271	1	1230	MRKTQCKNAESSKNQNASFPP
			į			KDHNSLTAREQNWIENEFGELT
1						EVGFRRLNQEEVEFLNRPITGSE
						IEAIINSLATKK\GPEGFIAEFY*R
						KTQCKNAESSKNQNASFPPKD
					:	HNSLTAREQNWIENEFGELTEV
						GFRRLNQEEVEFLNRPITGSEIE
				•		AIINSLATKKRTRGIHSRILLEVI
				ŀ		TSFYVERGGNAMSFMGKGVTK
						STILCLLHLSHEMMAQAGSLE
						WMSLWFLPLGSHSEEHIPTQQG
ĺ				•		LAWLIPLWVDRDPEVRFTSLGL
						GSALTTLETGCVALANSCQNIS
						GGLWGTVVNILLDQSECSMVR
		ļ				RESQHHVDVAKVYHLHPPEW
						WPKLYLGPLDHSRELRSSVLEC
						MEQRLEVTLGTTPYFSVPISKFI
						QAVTTKCLRLGASRFSVCLAGL
						SAAPEQKIHPPPPTQQFQQLPHD
						SVGFLQFSLHPKGNFGSGISSPD
						RSGDASHYSYPSLWQIRGTSRV
İ						KKQTWGSGNLKKWGIHKVQP
				1		QRLMMQNDTDEESI
21145	51513	Α	21272	1	987	QKEMMQNDTBEESI
21146	51514	A	21273	180	466	ARGSERAAASFSASMYEGKKT
[						KNMFLTRALEKILADKEVKKA
						HHSQLRKACR\VALEGNKAETE
						KQSPPHGEAKAGISTLPPVKSK
	1					TNFIEADKYF
21147	51515	Ā	21274	1	1224	
21148	51516	Α	21275	251	5986	SCRCPPRLQSPTSTPSPFSLLFLP
	ŀ					LLLLPSSRWEPRVGRAAASFSA
					Ì	SMYEGKKTKNMFLTRALEKIL
						ADKEVKKAHHSQLRKS/CGRV
						ALEEIKAETEKQSPPHG\EAKAG
						SSTLPPVKSKTNFIEADKYFLPF
						EL\ACQSKCP\RIV\STSLDCLQK
		:				L\IAYGAL*LGNAPDSTTPGKKL
		ł				IDRIIETICGCFQGPQTDEGVQL
						QIIKALLTAVTSQHIEIHEGTVL
						QAVRTCYNIYLASKNLINQTTA
						KATLTQ
21149	51517	A	21276	46	214	KLRLTTTSVYAIPNRRRYDGFK
				_		LVQSCIFI*NFRS*DLRPPQYMPF
		1				LIEGGMTVSNSCRAAFSFTSLC
				1		RHFRSEAWLQNTNTGGFGSTF
21150	51518	Α	21277	306	616	
21151	51519	В	21278	47	927	
		<u> </u>	<u> </u>			<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
21152	51520	A	21279	660	890	GGKITASGSPVPRPSGA*ALPSG
						RCSTPRTSSCPRQLPEPPGL\PSP
		l				PSAPANTTPDDSHAAP*GTRAH
						THRPPPHT/PTAPSRPPTSPPVAT
						RTAPKAPQ/HHP/PTSCPTQRPT
						DSPSVHEPSQPLATQHSTQPI*P
						VIVTQ
21153	51521	Α	21280	3	213	
21154	51522	A	21281	349	2616	TTAMAHRKLESVGSGMLDHRV
	1					RPGPVPHSQEPESEDMELPLEG
	1	1				YVPEGLELAALRPESPAPEEQE
						CHNHSPDGDSSSDYVNNTSEEE
						DYDEGLPEEEEGITYYIRYCPED
		1				DSYLEGMDCNGEEYLAHSAHP
						VDTDECQEAVEEWTDSAGPHP
		1				HGHEAEGSQDYPDRQLPIPEDE
						PSVLEAHDQEEDGHYCASKEG
						YQDYYPEEANGNTGASPYRLR
		1				RGDGDLEDQEEDIDQIVAEIKM
						SLSMTSITSASEASPEHGPEPGP
						EDSVEACPPIKASCSPSRHEARP
						KSLNLLPEAKHPGDPQRGFKPK
						TRTPEERLKWPHEQVCNGLEQP
						RKQQRSDLNGPVDNNNIPKTK
						KVASFPSLVAVPGPCEPKNLID
}						GIIFAANYLGST\QLLSERNPSK
						NIRMMQAQEAVSRVKRMQKA
						AKIKKKANSEGDAQTLTEVDLF
						ISTQRIKVLNADTQETMMDHAL
						RTISYIADIGNIVVLMARRRMPR
		1				SASQDCIETTPGAQEGKKQYK
						MICHVFESEDAQLIAQSIGQAFS
						VAYQEFLRANGINPEDLSQKEY
						SDIINTQEMYNDDLIHF\SNSEN
		1				CKELQLEKHKGEILGVVVVESG
		1				WGSILPTVILANMMNGGPAAR
		1				SG\KLSIGDQIMSINGTSLVGLPL
		1				ATCQGIIKGLK*PDHR*SFNIV\S
						CPPV/TPRFLYSKRPDLKYQLGF
						SVQNGIICSLMRGGIAERGGVR
	_ <u></u>			<u> </u>	<u>L</u>	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21155	51523	Α	21282	20	771	TTPNKQHMMQTRAGSAWYLR SRGCSQGAPQKASSGSGGSSW NRSVATVPGIVLGWRRPELLSE RGCAEARGAGVAGRQRLQLTG VQGKSVVRGRGLLGRREWCRR GGGAAPRPTPAGTPVPRRPGTR AGPEEGEGAGAGARAAPEPRP HALSPPAAFGADGQLRPLLLTS LRAHHPGRGGLLLTSPGTHQPR RGRQGTPPPASPSPA/PEEGALP GVLPL*PPPSPPSP/SSESPRPHPG GGALAVTRRTAAA
21156	51524	Α	21283	1615	1985	LGSTRQAAPGQLRGHQAPAPN SDATAGQGRCRRSAD*GWGSR EQI*ALPLPSGPAPLLPPLLPPLL PVPPSPPPPSPPSPPSPLPPP\GPP* PPLTFSVVTPGGLWCPPHRKSL NPTSKGVNEY
21157	51525	Α	21284	3	501	
21158	51526	Α	21285	1	2646	
21159	51527	A	21286	31	319	VIQDLHVPPREESAEQVDDPPE PVY\RT*RGSPGPLHRAPLQPPF PARCGRRTRTRAPGAPTTTTQT RELPPGSRPLRLPRVPPAQPPPL PRWTAT
21160	51528	Α	21287	3	147	
21161	51529	A	21288	20	600	PVTPRRTILPWALSVSPALPLP* PHPPAGLVMSARTSRCSTPTTS LRSSG*GWRTPTGSHTSTIQRTP LFDGSCPRSLSLPLEASINPARM VTPQPRPALQRRRSQQSWMRL GAGRKSLLPQLL*GPRPWTRQG CSIAPRRQTRESGSGRSTGVPPG LCWRVAS*HSSRTQRPRLQAA* GSLPSFPPLSTQWS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21162	51530	A	21289	383	1800	TCSTQTSLRNPPRRPRAASLPQT
						CSRSPRPSRAHSLTGRPPRFQSQ
						MPSRSPCC*IC*NIRIDLCGS\TH
					ľ	SDGRFELVFSGKKLALRASSQD
						EAEDWLDRVREALQKVRPQQE
					ļ	DEWVNVQYPDQPEEPPEAPQG
						CLSPSDLLSEPAALQGTQFDWS
						SAQVPEPDAIKESLLYLYMDRT
						WMPYIFSLSLEALKCFRIKNNE
					ļ	KMLSDSHGVETIRDILPDTSLG
						GPSFFKIITAKAVLKLQAGNAE
İ						EAALWRDLVRKVLASYLETAE
						EAVTLGGSLDENCQEEKNNTK
						HTTARNAFEENDFMENTNMPE
1						GTISENTNYNHPPEADSAGTAF
						NLRPTVKQTETKWEYNNVGTD
						LSPEPKSFNYPLLSSPGDQFEIQ
}		1				LTQQLQSLIPNNNVRKLIAHVIR
		1				TLKMDCSGTHVQVTCAKLISRT
1		1				GHLMKLLSGQQEVKASKIEWD
		1				TDQWKIENYINESTEAQSEQKE
		1				KSLEAASGTQDGVHVQEPRPQ
1		1				APSPLDLQQPVESTSSQQPSSTV
		1				SETAREVGQGNGLQKAQAHDG
						AGLKLVVSSPTSPQQVCKLWW
						SAGLHTPPPVTSLGTVGLQGPQ
						QRFPKFPDFFHPLEISCGKVA
21163	51531	A	21290	2	667	
21164	51532	Α	21291	84	377	RMGKHCFENLLFRDGKAQRAC
		1	ŀ			MGYTVCHSHIVGSGQKCWPA
		1				WPQVNLQPSRHQQVCCSLPRC
						AKCKTVFHQSCQAV\VKKGCPR
		_				CAR*RKYQQQNVFA
21165	51533	Α	21292	2	1168	FVLPVVSGRETASSARRLRGCR
		1				SREPPSPRSSLRLVSCDSLREQL
						REWRCFQWWRMDWTPRLPSR
		1				HIISELEHLTFVNMDVGRCRAW
		1				LRLALNNGLMECYLKLLLQ\EQ
		1				ARLREYYQPTALLRDAEEGEFL
		1				LSFLQGLMSLSFELSYKSAILNE
	t .					WTLTPLALSGLCPLSELDPLSTS
						GAELQRKESLDSISHSSGSEDIE
			]		1	VHHSGHKIRRNQKPTASSLSLD
1					]	TASSSQLSCSLNSDSCLLQENGS
						KSPDHCEEPMSYDSDLGTANA
		1				EDSDQSLQEVLLEFSKAQAASG
		1				TQDGVHVQEPHPQAPSPLDLQ
İ		1				QPVESTSGQQPSSTVSETAREV
		1				GQGNGLQKAQAHDGAGLKLV
		1				VSSPTSPNISSMIQSTRPVHFCIV
	<u> </u>	<u>L.</u>				STKHCTWEKVGMDE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21166	51534	A	21293	263	813	
21167	51535	A	21294	130	1094	NOPHRTKGCPNSSTECKNGSLT
21107	31333	ľ`	21254	130		CDFMQVPILEMGKLRPEGERRV
						QDRDSNPSLSPSAGLFQARWS*
		1				PPRPSFLLPAGKRRGCDGGGRG
						GGRVRAGGAPLRVHRQGRAPP
						QSTEHWWHVRRKPGGRPFYLP
						AQYVRELPALGNPAADAPPGPP
				•		IGTRSP*AARLRLPVRERGCGSG
						PQRPPSGAPRRGQLPVRPCAAR
						RREP/RAAAWRPACPPACTCGP\
						GARAARAVPGRPDARRRLASR
						RQPRKQRQLQGLQRSGLLGVP
				1		AAPVAQRLRERLRGHPGGTRP
						AAVGDREAGTTPGAEAGRLAE
						DSARREPGLEENHSAQRRA
21168	51536	A	21295	2	3676	
21169	51537	Α	21296	432	3453	DLAGEVLLCGGAADYGTLGFS
						SPAVQEANRGALCSSIERGHIIS
						ELEHLTFVNTDVGRCRAWLRL
						ALNDGLMECYLKLLLQEQARL
						HEYYQPTALLRDAEEGEFLLSF
					į	LQGLTSLSFELSYKSAILNEWTL
						TPLALSGLCPLSELDPLSTSGAE
					· ·	LQRKESLDSISHSSGSEDIEVHH
						SGHKIRRNQKLTASSLSLDTASS
						SQLSCSLNSDSCLLQENGSKSP
		ł				DHCEEPMSCDSDLGTANAEDS
						DRSLQEVLLEFS
21170	51538	Α	21297	64	2996	LCPPAQPAASQPQVELPAGWPQ
						PHALPLPHDHRAQHLLGCAHP
		İ				GRAKPELQQHHDCACAAQIPHI
						PVRQPYQHPDPPLCQPRRPACP
						ALPIHGRQLLLQEPLQAGTGGG
						PGCPPWPGQPHPPVTQVQQPHC
						GAPQPAFQPGVSAVVLRGRIVK
						LAPEDLANLTALRVLDVGGNC
	1			!		RRCDHAPNPCMECPRHFPQLHP
	1					DTFSHLSRLEGLVLKDSSLSWL
						NASWFRGLGNLRVLDLSENFL
						YKCITKTKAFQGLTQL
21171	51539	A	21298	1	320	PVRTPPIPAFSPLKTLRIMSLHQF
- 1 / 1	31339	1,	21270	1	320	LLEPNCGRPQLCRGTMARKPAP
						RKH*KHISRQCRVVLFEYKIGES
						-
						VGFFKAVIFLFVFLRFHSILDQS
21172	61640	_	21200	_	142	FSLSSLLWKIVTLT
21172	51540	Α	21299	1	1050	
21173	51541	Α	21300	1	1059	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21174	51542	A	21301	116	1559	PRTARPSCAHARTVRAVATRRR RVHFRDLSAAPSGLSVLRPRPT EPVRVDCPESANPPVRARPDSP DPRMFELLAASDARINGPFCAP RVLLAPLLGAFVTVSKIVSKIAL SPK*SRRSHLLRRNGSQW\VKV H\ELKEHNG\HITGID\WAPKSD RICH/CLGADRNCLMVWS\QK\D GVWKPNPG*FLRNLIGGGNFL* KWVPPRRTKFGCGGSGARLISV CYFESENDWWVSKHIKKPIRST VLSLDWHPNNVLLAAGSCDFK CRVFSAYIKEVDEKPASTPWGS KMPFGQ\LMSEFGG\SGTG\GW VHGVSFS\ASG\DRLGWGSATD ST\VSVA*CLQKVVQVSTLKT\E FLAAP*VVSFVSREPASGLPAH DC\CPMLFNYG*PRAALTFRLP KLDIPK\QSIQRNMSAHGNAFG NM\DKRATTED\RNTAL\ETLH\ QNSITQVLYLLRWDKQD\CRKF CT\TGI\DGEPWTIWDFQDPRSL PSOGLRIM

2695

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21175	51543	Α	21302	1	2223	MKHYTRVQSSISPPAERQETHA
						WASPAVTSLESAACHELQEAD
						LSESLSYPRIVSSSSLQQYVAQG
						GSFPCFGMPWNFISGGAESTNA
ļ						VISFANATTAVPMAVLSRRESS
						LANNPGVVNYSALPENENVGP
						EMPERPANSSKNSTETANYPTL
ļ						MGNYNGQNTASLSVFIPPYFAE
		1				KIILTEMPGTTETNVENNSQTV
						YYPALSGNTSAPYPASSYLPITS
						NFVLERVAEMPETAGTNVKNN
						TQTVNYPPSSGRGRSQGNASLS
ŀ						LSIPPKFGIERHLGCFLVLAIRSK
						AAMN\IIYHIGFV*TQAFISLRY/
						MPKSAVAGSYGNALQNVLHIE
						MLFKTESGPQMSYGTMSYSTE
						MKNNCDQDDASASACLTPDFA
						LLPLNILVKVDTNTENSVNTMN
						RSTLLDSDSGQDSSSSSVCIPPK
						YGYLGDPKRNVRVLKIHLLAV
						QNMAKPKQAACYLVRILFSKEI
						LISSSVDIHLKDSQSLDPNKMA
						ALREYLATTFPTCDLHEHGKD
i	1					WQDCISGINSMIYCLCSEGKSTP
1						KTVRKNKKRTNRVASASADRN
						DQRGRDGGEGCSWMFQPMNN
						SKMREKRNLQPNSNAIPEGMRE
		ŀ				PSTDNPEEPGEAWSYFGRPWRN
					Ì	IRMPCSVLTLAKTKSCASLSAR
						YLIQKLFTKDVLVQSNVYGNL
						KHGLCALDPNKISALREFLQEN
						YPICDLSENGRDWKSCVTSINS
21176	51544	Α	21303	1	297	
21177	51545	Α	21304	1	162	
21178	51546	Α	21305	11	212	KGKNCVAIAADRRFGIQAQMV
						TTDFQKIFP/MGAQRLKFRLNL
						YELKEGRQIKPYTLMSMVANL
						LYEK
21179	51547	Α	21306	37	554	SSIKFGRSNPDYWIALPSRNLDP
						AIMSIMSYNGGAVMAMKGKN
					1	CVAIAADRRFGIQAQMVTRDF*
						KFFPMG*PACTPGMAGFAIDVQ
					1	TVAQRLKFRLNLYELKEGRQIK
						PYTLMSMVANLLYEK/RDPDHL
						FETISQAMLNAVDRDAVSGMG
						VIVHIIEKDKITTRTLKARMD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide sequence	1	in USSN 09/540,217	location of first codon for peptide sequence		*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21180	51548	A	21307	51	741	YTAIMSIMSYNGGAVMAM\KG KNCVAIAADRRFGIQ\ARLLTTN FQKIFPMGDRLYIGLGGASPLD VQTS/VAQRLKFRAEPVMS*RE GRQI\KPYTLH*AMGGQPSLL*E KVFGPLTYTE\PVHLPGLGPERP FKALSICFSKTLIRVGPMGDLN DFCGSSGNLAPNKMYG\MCESL WGGPTWDPDHLF*KPFSQGHG
						*MLWDRDGS/VSGMGSSLFHIIE KDK\IPPRTLKARMD
21181	51549	В	21308	150	404	
21182	51550	Α	21309	199	395	
21183	51551	A	21310	100	182	SFLFWIKR*SFPDALYFSSKSGA CSCN
21184	51552	A	21311	203	289	KCLALDIALATRRSAI*FPSVKG PGRGV
21185	51553	В	21312	166	411	
21186	51554	A	21313	232	317	KCPALDIAVATRRSAI*FPAVKG PGRGV
21187	51555	A	21314	129	278	FYSSVSGVSSKDDRRFGCSGYS AAVISFRQC*SQLPGCSATYQR ESTFS
21188	51556	Α	21315	789	962	
21189	51557	A	21316	477	599	
21190	51558	A	21317	86	577	ATEPLAPSERREVSIPLLVPAPL PPPRPPVPSRTRTRPWRRP*AC FPMTLAASCGPTRSPWPSQQPIP QPAPVG/PGNIKTLGDAYEFAV DVRDFSPEDIIVTTSNNHIEVRA EKLAADGTVMNTFAHKCQLPE DVDPTAV\TSALRKDGSLTIRAR RDPHTEHV
21191	51559	В	21318	312	466	
21192	51560	A	21319	886	1338	GFWQVIRALRGGEVHREGTSA PGRHRRRCGCRSAWNRCCTAC SHVWPPRQMWPQG/PDVAMKT GNIKTLGDAYEFAVDVRDFSPE DIIVTTSNNHIEVRA/REA/ERPD GTVMNTFAHKCQLPEDVDPTS VTSALREDGSLTIRAWRHPHTE HVQ
21193	51561	A	21320	3	605	KKLTPVVRSRVERDPRVRRGLR PHPVLSSLRPPGQGPAPSVDEPQ NLFHLPSGEKIPFLFLLLLFTPP PRPPVLSRPRTRPWRRP*ACFP MTLAAICGPTRSPWPSQPAPV/V AGNIKTL*DAYEFVVDVRDFSP EDIIVTTSNNHIEVRPEKLAADG TVMNTFAQQCQLPEDVDPTSV TSALREDGSLTIRARGHPHTEH

SEQ ID	SEQ ID NO:		SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence	İ	09/340,217	sequence	or peptide sequence	detection, (-possible nucleotide insertion)
21194	51562	Α	21321	3	450	SSAAEVQEPRGAPGSPSIPSASR
•						PGPAPRPASLPPSQMQAAALPE
						EIRWLLEDAEEFLAEGLRNENL
						SAVARDHRDHILRGFQQIKA\G
						GDIGQDSSDDNHSGTLGLSLTY
						DAPFLSDYQDEGMEDIVKGAQ
						ELDNVIKQGYLEKKS*DH
21195	51563	A	21322	1	399	IGQDSSDDNHSGTLGLSLTSDA
						PFLSDYQD/EG*VHQISDHWEK
						LRTWC*VYCDKGMEDIVKGAQ
						ELDNVIKQGYLEKKSKDHSFFG
						SEWQKRWCVVSRGLFYYYANE
						KSKQPKGTFLIKGYSVRMAPHL
						RRDS
21196	51564	Α	21323	3	1189	RLPARPPPDQGPRPVPPLSRPAK
						CRLAVLPEEIRWLLEDAEEFLA
						EGLRNENLSAVARDHRDHILRG
						FQQIKARYYWDFQPQGGDIGQ
				:		DSSDDNHSGTLGLSLTSDAPFL
						SDYHDEGMEDIV*GAQELDNVI
						KQGYLEKKSKDHSFFGSEWQK
						RWCVVSRGLFYYYANEKSKQP
						KGTFLIKGYGVRMAPHLRRDS
						KKESCFELTSQDRRSYEFTATSP
	}					AEARDWVDQISFLLKDLSSLTIP
						YEEDEEEEKEETYDDIDGFDS
						PSCGSQCRPTILPGSVGIKEPTEE
						KEEEDIYEVLPDEEHDLEEDES
						GTRRKGVDYASYYQG\LWDC\
						YGDQPDDLSL\QRGDLILFLSKE
						YNMYGWWVGELNSLVG\IVPK
						EYLHHLPLKWKERWKPREIYFF
21197	51565	Α	21324	47	225	NSHHVRGRPRCADSSSPSGDRG
						QPEAQPAPDSSAPEHAQEPGRA
						AVKRPDL*SHMTRRP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
21198	51566	A	21325	66	725	AILIILLSSEGLWSSDQHRLVGV
						QDSPPQGSLGCYFSAMATSEQS
						VCQARASVMVYDDTSKKWVPI
						KPG\QQGFSRINIYHNTVSNTFR
						VVGVKLQDQQ\VVINYSIVKGL
						KYNQATPTFHQWRDARQVYG\
						LNFASKEEATTFSNAMLFALNI
1						MNSQEGGPSSQRQVQNGPSPD
						EMDIQRRQVMEQHQQQRQEFL
<u> </u>						ERRTSATGPILPPGHPSSAASAP
						VSCSGPPPPPPPPVPPPTGA\TP
						PPPPPL\PAGGAQGSSHDESFHV
İ						QDWPLAISWGPSLRRVPNGPED
						ASGGSSPSGTSKSDANRASSGG
						GGGGLMEEMNKLLAKRRKAA
						SQSDKPAEKKEDESQMEDPSTS
						PSPGTRAASQPPNSSK\AGRKP
						WERSNSRGYFCVLDSVQNPAL
-						WPKSPES*EPPSVSSL/ISRMKPA
						GS\VSDMA\LDAFDLD\RMKQEI
						LEEVVRELHKGERKEIID\AIRQ
						EA*SGISRKKNLGHRAHPPTRTS
						FICSQRPRLM
21199	51567	Α	21326	301	390	
21200	51568	Α	21327	72	415	
21201	51569	Α	21328	395	697	RVMSPVHVNTICSPHPHSDGDV
						IVPIVLSNERYHWP\VLGWVGG
						VDCDEFFRVVLGEARLH\LDGV
						AHQH*TKEGTISNLFQ\ATSVSH
						LVEVVPQKLSWVI
21202	51570	Α	21329	36	399	VEILTSKRAGDWFKEELRLLGG
						LDHIVDKVKECVDHLSRDEDEE
						KLVASLWGAQRCLRVLQSVTV
						HNPENQSYLIAYKDSQLIVSSA
						KALQHCEELI*QYNRAVDTLCL
	<u> </u>	<u> </u>	21225		100	ADCKPLPHPTVL
21203	51571	A	21330	1	177	
21204	51572	C	21331	198	350	
21205	51573	A	21332	113	228	
21206	51574	Α	21333	1	466	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	i .	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21207	61676	<del>                                     </del>	21224	<u> </u>	4122	L ASGCWRKEVGAGSANGVEMV
21207	51575	Α	21334		4133	!
						QGPVQTPALTIHRRKRRRRRR
		1				RPSGGTQQALLPAPAVAASES*
		1		1		ARARPAALEAGREVAGPP\SCPP
		İ				NRASYLPKAESLASLGSHLPAL
						LSRARVPRPPAGRKERE\RRKRP
						\VAKAPARL\RGEYETG\VKMTS
						RFAKTYSRKGGNGSSKFDEVFS
					1	NKRTTLSTKWGETTFMAKLGQ
						KRPNFKPDIQEIPKKPKVEEEST
						GDPFGFDSDDESLPVSSKNLAQ
						VKCSSYSESSEA
21208	51576	Α	21335	3	221	
21209	51577	Α	21336	1	580	
21210	51578	A	21337	2	1095	ARGNRPPPPPRTERGVLESRTRS
		ä				RHRAPRLPPAMPAPRAPRALAA
						AA\PASGKAKLTHPGKAILAGG
						LRGG\IQICMTFPTEYVKTQLQL
						\DERSHPPRYRGIGDFVRQTVRS
						HGV\LGLYRGLSSLLYGSIPKAA
				1		V\RFGMFE\FLSNHMRDA\QGRL
						DSTRGLLC\SLGAGRGPRP*VV
		1				VCPMETIKVKFIHDQTSPNPKY
						RGFFHGVREIVREQGL\KGTYQ
		1				GLTAT\VLKQGSNQA\IRF\FVM
						TSLRQW\YPRGTTPNKAHEPF*I
						TWGSFGA\IASASQVSFGNTPLD
1						VI\KTRMPGPWRPHK\YR\NTWD
						\CGL\QILKKGGGSRAF\YKGTV
						PPLGPGLAWDVAI\VFV\IYDEV
		1				G*SCFNKVWKNGIKA
21211	51579	A	21338	3	375	TAGDSGAEDGPASRVSQMPTLP
						PPMPWNLPAGVDCTTSGVLAL
						TALFFKMEEANLASRAKAQELI
						QATNQILSHRKPPSSLGMTPAP
						VPTSLGLPPGPSSYLLSGSPPLG
						GC\GSTPPTPPGLAA
21212	51580	Α	21339	51	3789	GRGHPHGGGELLPASPRGSPEG
					1	DQPGQGDPEEKEGRVVPRGGA
				1		PDVPSAHALVPEPQTVLGLHQR
				1		VTAQPGRHGAH*LHGDRLHSG
		-			1	RPQRPTHTGWAPCLPYPTLGSG
						*RRHHGNPAGTPRSLLFPF\PEK
						ELEAPPPAPRFDIYDPFHPTDEA
						YSPPPAPEQKYDPFEPTGSNPSS
						SAGTPSPEEEEEEEEEEEE
						EEEEGLSQSISRISETLAGIYDD
						NSLSQDFPGDESPRPDAQPTQP
				-		TPAPGTPPQ
21213	51581	A	21340	286	393	
21213	51582	A	21341	64	610	
21214	21202		21341	I <sup>V¬</sup>	1010	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
21215	51583	Α	21342	1	4433	MWAQLLLGMLALSPAIAEELP
						NYLVTLPARLNFPSVQKVCLDL
!						SPGYSDVKFTVTLETKDKTQKL
						LEYSGLKKRHLHCISFLVPPPAG
						GTEEVATIRVSGVGNNISFEEK
						KKVLIQRQGNGTFVQTDKPLYT
		1				PGQQVYFRIVTMDSNFVPVND
1		ł			ŀ	KYSMVELQDPNSNRIAQWLEV
						VPEQGIVDLSFQLAPEAMLGTY
					-	TVAVAEGKTFGTFSVEEYVLSP
	-					FLLLLSSVLPKFKVEVVEPKELS
						TVQESFLVKGGL*\
21216	51584	Α	21343	3	522	ELRTWIEGLTGLSVGPDFQKGL
						KDGTILCTLMNKLQPGSVPKIN
					:	RSMQNWHQLENLSNFIKAMVS
						YGMNPVDLFEANDLFESGNMT
						QVQVSLLALAGKMGTNKCASQ
1						SGMTAYGTRRHLYDPKNHILPP
						MDHSTI\SLQMGTNKCASQVG
						MTAPGTRRHIYDTKLGTD\KCD
21217	51585	Α	21344	1	1230	MHQEDLRAWYLDLGLPSHQNE
						TTDVSVVYERFSLTDATYPKQA
						KQWDFPFLRVSTAQPTAWKCQ
		ŀ				RAPSPYTHQDTALIPSPTARWL
1						SPEKEPKQGEVGEKSLLPDPTLP
						LTDPRLTGSTEQKKDLAAGPVG
						SALRRRPPAVSSTQFNKGPSYR
1	Ì					LLADVQNRLLPKYDSQKEAEL
						RSWIKGFTGLSIRPDFQKGLKD
1						GIILCTLVNKLQPGSVPKIN\ASV
						*NWH*LENLSNFLKAMVSYGM
						/NPVDLFEANDLFESGNNMQVR
ļ						VSLLALAGKAKTKGLRSGVDIR
						DKYSKKQNFDDTTMKASQCVI
}						RLQITNKCASQSGMTAYGTRR
}						HLYDPKNRILPPMDNSTISLQM
						GTNKYASQVGMTAPGTQRHIY
						DTNLGIDKCENSSMSLKMGYT
		<u> </u>				QVANHSGQVFGLGRQIYEPKY
21218	51586	Α	21345	2	389	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21219	51587	Α	21346	1	1113	MHQEDLRAWYLDLGLPSHQN
						AQPTAWKCQRAPSPYTHQDMA
						LIPSPTARWLSPEKEPKQGEVG
						EKSLLPDPTLPLTDPRLTGSTEQ
						AHAEGLAALMSALRVSHLQGR
						GGVVTLVDSQLGVIAVSSTQFN
						KGPSYRLLADVQNRLLPKYDS
						QKEAELRSWIKGFTGLSIRPDFQ
İ		ŀ				KGLKDGIILCTLVNKLQPGSVP
		1				KIN\ASV*NWH*LENLSNFLKA
						MVSYGM/NPVDLFEANNLFESG
						NNMQVRVSLLALAGKAKTKGL
						RSGVDIRDKYSEKQNFDDTTM
						KASQCVIRLQITNKCASQSGMT
						AYGTRRHLYDPKNRILPPMDNS
						TISLQMGTNKCASQVGMTAPG
						TQRHIYDTKLGIDKCENSSMSL
21220	51588	A	21347	353	1260	SCWCLKLALAVSST/QFNKGPS
21220	31366	Γ	21347		1200	YRLLADVQNRLLFKYDSQKEA
						ELRSWIKGFTGLSIRPDFQKGLK
						DGIILCTLVNKLQPGSVPKIN\AS
						V*NWHQLENLSNILKAMVSYG
		-				M/NPVDLFEANDLFESGNNMQ
						VRVSLLALAGKAKTKGLQSGV
					}	DIRDKYSEKQNFNDTTMKARL
						CVIRLQITNKCASQSGMTAYVT
						RRHLYDPKNRILPPMDNSTISLR
						i i
						MGTNKCASQVGMTAPGNQWH
						IYDTKLGIDKCENSSMSLKMGY
						TQVANHSRQVFGLGRQIYEPKY
0.1001	51500	<u> </u>	21249	2	520	QPGGPVAHGAPSAGNCPGPGE
21221	51589	A	21348	3	530	MPESWLAAFPPYGDIKKVVGL
21222	51590	Α	21349	1	853	QLKEGLHQNLTMQDSIQIFDFQ
						LPGLLRPSSVLWTLKPYHTAGL
		1				ASDRDSYKGILACMPCMFGSRR
						SPLRDIKSHQADLELEVKNSLD
				[		TIHRLESELKKQSKIQSQMKVE
				1		KAHLEEEIAELKKSQAQDKAKL
				1		LEMQESIKDLSAIRADLANKLA
						EEERAKKTSA*GAFGP/LIAQAN
						SRDEGTATIITQLELERDVHQRE
						LKDLTSSLQSVKTKHEQNIQEL
				1		MKHFKKEKSEAENHIRTLKVPL
					L	LNYPDDTEGIFLVAHEVNKA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21223	51591	A	21350	1	652	NTSLINTKKKLETDISQIQGEME
						DIIQEARNAEEKAKKAITDAAM
ļ						MAEELKKEQDTSAHLERMKKN
						LEQTVKDLQHRLDEAEQLALK
						GGKKQIQKLEARVRELEGEVES
						EQKRNVEAVKGLRKHERKVKE
İ						LTYQTEEDRKNILRL\QDL\VDK
						LQAKVKSYKRQAEEAEEQSNV
			9			NLSKFRRIQHELEEADF\RADIA
						ESQVNKLRVKSREVHTKIISEE
21224	51592	Ā	21351	21	5868	PAAMSSDSEMAIFGEAAPFLRK
						SERERIEAQNKPFDAKTSVFVV
						DPKESFVKATVQSREGGKVTA
						KTEAGATVTVKDDQVFPMNPP
						KYDKIEDMAMMTHLHEPAVLY
						NLKERYAAWMIYTYSGLFCVT
						VNPYKWLPVYNAEVVTAYRG
						KKRQEAPPHIFSISDNAYQFML
						TDRENQSILITGESGAGKTVNT
						KRVIQYFATIAVTGEKKKEEPA
						SGKMQGTLEDQIISANPLLEAF
						GNAKTVRNDNSSRFGKFIR
21225	51593	A	21352	54	399	
21226	51594	A	21353	441	1585	VPPMGGGPPFVGPVGFGPGDRS
						HLDSPEAREAMFLRRAAVAPQ
						RAPILRPAFVPHVLQRADSALSS
						AAAGPRPMALRPPHQALVGPP
						LPGPPGPPMMLPPMARAPGPPL
						GSMAALRPPLEEPAAPRELGLG
						LGLGLKEKEEAVVAAAAGLEE
						ASAAVAVGAGGAPAGLAVIGP\
						SLPLALAMPLPEPEPLPLEVV
						RGLLPP\LRITELLSLRPRPRPPR\
						PEPPPGLMALEVPEPLGEDKKK
						GKPGEIEHRCIATA/AQGASWE
						DPPTCWKWDAE*LSRVFCGDL
						G\NEVND*HLGHAASARFPIPSL
						KAKVIR*QSAQARPKGYG\FVS
						FKDP\SDYVRP\MREMNGKYVV
						LAPPIKLRKSMWKDRNLDVFR
	<u> </u>	L.,				KKQKEKKKLGLR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	,	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21227	51595	A	21354	1	1122	MPGGKIGFLIEKGATEQSSTGN
						DFDELREEGGRRSNYSELQEDI
						QTKGKEVENFEKNLEECVTTIS
	<b>]</b>					NTEKCVKELMELKTKARELRE
						ECRSFRSQCNQLEERVPAMKDE
						MNEMKREGKFREKRIKRNEQS
						LQEIWDYVKRPNLPLIGVPESD
						RENGTKLENTLRDIIQENFPSLT
						RRGQPFRIMENTENAPKDNLLE
						KEQLPRHNNCLELTQEEPAWD
	,					LQFDLLDLNDISSALKHHEEKN
						LISLLPQRGADYLSFRSHFQKNF
						VCVENCSLQERTVTGTVKVKN
						VSFEKKVQIRITFDSWKNYTDV
						DCVYMKNVYGGTDSDTFSFAI
						GLPPVIPTEQKIEFCISYHANG\Q
						IF\WDNK*WGRKYKNCVIVQW
21228	51596	Α	21355	71	1124	GPPPLCLMSCTRMIQVLDPRPL
						TSSVMPVDVAMRLCLAHSPPV
	ļ .					KSFLGPYDEFQRRHFVNKLKPL
	}					KSCLNIKHKAKSQNDWKCSHN
						QAKKRVV\FADSKGLSLTAIHV
İ						FSDLPEQPAWDLQFDLLDLNDI
						SS\ALKHH\EEKNLILDFP\QPST
						DYL\SFRSHFQK\NFVCL\ENCSL
						Q\ERTVTGTV\KVKNVEF*GRKF
						QIRYHFSDSWEKLTLDVDCVY
						MENVY/GVGTGSDTFSFAIDLPP
						VIPTEQKIEFCISYHANGQVFW
						DNNDGQ\NYWIVHVQWKPDG
						VQTQMAPQ\DCAFHQTSPKTEL
						ESTIFGSPRLASGLFPEWQSWG
						RMENLRLLCDELSNNVTGLDLS
						YSPMQS
21229	51597	C	21356	204	273	
21230	51598	В	21357	1	1062	OP DI* PTV DICODIVOV CHOPITO
21231	51599	Α	21358	1	397	QRPI*PTKPIGQRVSKGHQP\TR
İ						H\DFTQF\LAAHETISRLVGFSA
						GQV\QYLDLIKK\DPSKLFNEER
						LIDKTKVTYLKWLPESESLFLA
						SHASGHLYLYNVSHPCASAPPQ
	<u> </u>					YSLLKQGEGFSVYAAKYNDLF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
ĺ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21232	51600	Α	21359	71	1736	FRTSLKDVWHTPLTPFLPPLKSI
						DLNKPIDKRIYKGTQPTCHDFN
						QFTAATETISLLFGSFSPRRTWA
						WCGPAV*AGPCPTAFLGTLRLI
						DKTKVTYLKWLPESESLFLASH
						ASGHLYLYNVSHPCASAPPQYS
						LLKQ\AWGFSFYAAKSKAPRNP
						LAKWAVGEGPLNEFAFSPDGR
						HLACVSQDGCLRVFHFDSMLL
		İ				RGLMKSYFGGLLCVCWSPDGR
						YVVTGGEDDLVTVWSFTEGRV
						VARGHGHKSWVNAVAFDSLY
						TTRAEEAATAAGADGERSGEE
						EEEEPEAAGTGSAGGAPLSPLP
		1	}			KAGSITYRFGSAGQDTQFCLW
						DLTEDVLYPHPPLARTRTLPGT
						PGTTPPAASSSRGGEPGPGPLPR
						SLSRSNSLPHPAGGGKAGGPGV
						AAEPGTPFSIGRFATLTLQERRD
						RGAEKEHKRYHSLGNISRGGSG
						GSGSGGEKPSGPVPRSRLDPAK
						VLGTALCPRIHEVPLLEPLVCK
				į		KIAQERLTVLLFLEDCIITACQE
						GLICTWPRPGKAFTDEETEAQT
						GEGSWPRSPSKSVVEGISSQPG
21233	51601	Α	21360	2	467	LMQ/NWDAAMEDLTRLKETID
						NNSVSSPLQSLQQRTWLIHWSL
						FVFFNHPKGRDNIIDLFLYQPH
						MLADKLNMTPEEAERWIVNLI
						RNARLDAKIDSKLGHVVMGNN
				·		AVSPYQQVIEKTKSLSFRSQML
				ł		AMNIEKKLNQNSRSEAPNWAT
						QDSGFY
21234	51602	В	21361	1	909	
21235	51603	Α	21362	3	3617	YWKERPTQKVIPRATENHGLKS
						YLQKTKLSIDEAAFLLPDTNLK
						SELLELLTHWLQVGVPMTPSLG
						SINLLGWLTELRETHTYICWFIV
						KETTRDTDEEMCRTEPALACSI
						SHYCDDGCIQMLNTPETLQCSA
						KDSKHFIPKECSIPGENRPPSDT
						GKTVKFLSLNIFNLQLAESTDA
[						EQRANCILRCFLTETTLNYQKIL
						SVRPGTKLATASHVSGLGLQTP
						PFGLAQHLIRPHAFLAPKDPLTS
	<u></u>			_		FTERNSRSG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21236	61604	<u> </u>	01262	75	2141	GPRLGLTTMWHEARKLGRRLR
21230	51604	A	21363	75	2141	
						GLLFDSRKRAERRREYYEKIKK
						DPAQFLQVHGRACKVH\LDSA
			1			VALAAESPVNMMPWQGDTNN
						MIDRFDVRAHL\DHIPDYTPPLA
						HHHFSQNQESDER\KCNYERYR
						GLVQNDFAGISEEQCLYQIYIDE
						LYGGLQRPSEDEKKKLAEKKA
						SIGYTYEDSTVAEVEKAAEKPE
						EEESAAEEESNSDEDEVIPDIDV
						EVDVDELNQEQVADLNKQATT
						YGMADGDFVRMLRKDKEEAE
						AIKHAKALEEEKAMYSGRRSR
						RQRREFREKRLRGRKISPPSYAR
						RDSPTYDPYKRSPSESSSESRSR
						SRSPTPGREEKITFITSFGGSDEE
						AAAAAAAAASGVTTGKPPAP
						PQPGGPAPGRNASARRRSSSSSS
						SSSASRTSSSRSSSRSSRSRRGG
						GYYRSGRHARSRSRSWSRSRSR
						SRRYSRSRSRGRRHSGGGSRDG
						HRYSRSPARRGGYGPRRRSRSR
						SHSGDRYRRGGRGLRHHSSSRS
						RSSWSLSPSRSRSLTRSRSHSPSP
						SQSRSRSRSRSQSPSPSPAREKL
						TRPAASPAVGEKLKKTEPA\AG
						KETGAAKV\PKLTPQEKLKLRM
						QKPL\NRQFKADKKAAQRKR*S
						QAEHERQERKDEA\RAMARKIG
						MKERER\REKEREEWERQYSRQ
						SRSPLP/RDYSREYSSSRRRSRSR
1						SRSPHYRH
21237	51605	A	21364	1	609	JOKOT IT TOTAL
21238	51606	A	21365	1	1227	
21239	51607	A	21366	1	1626	
21240	51608	A	21367	370	4388	GFIWNRITPTEDGQTRELQNHL
2.270		<b> </b>	[	"	1.230	GLSAPVPRGRKGKKVKTQTSSF
						DIQKAEWLRKYNPEQLLQDEG
						YKKHIKHHCNKVLLRVRMLYY
						LKQEVIGNECQKVFDGVDASDI
						DVWVPEPDHSEVPAEWWDFD
						ADKSLLIGVFKHGYEKYNTIRA
						DPALCFLERVGKPDEKAVAAE
						QRANDYMDGDVEDPEYKPAPA
						IFKDDIEDDVSSPGDLVIADGDG
						QLMEGDKVYWPTQSALTTRLR
		<u> </u>		l	<u> </u>	RLITAYQRTNKNRQIQQIQ

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21241	51609	A	21368		530	AEADAIIQMVREGQRARRQQQ AATSESSQSEASVRREESPMDV DQPSPSAQDTQSIASDGTPQG\E KEKEERPPELPLLSEQL\ALDEL WDMLGECLKELEESHDQHAVL ETHRTVLNQILRQSTTHLADGP FAVLVDYIRVLDFDVKRKYFR QELERLDEGLRKEDMAVHVRR DHVF
21242	51610	Α	21369	1	217	
21243	51611	A	21370	274	599	PVSKESRVAPLCDFCLPFIQSES SQSEASVRREESPMDVDQPSPS AQDTQSIASDGTPQGEKEKEER PPELP/LLSEQLSLDELWDMLGE CLKELEESHDQHAVLVLQPA
21244	51612	A	21371	2	731	TPTPPTAPTPVTSAPALVAATAI STIVVAASTTVTTPTTATTTVSI SPTTKGSKSPAKVSDGGSSSTD FKMVSSGLTENQLQLSVEVLTS HSCSEEGLEDAANVLLQLSRGD SGTRDTVLKLLLNGARHLGYT LCKQIGTLLAELREYNLEQQRR AQCETLSPDGLPEEQPQTTKLK GKMQSRF\SGLGSASSIQAAVR QLEAEADAIIQMSESSQSEASVR REESPMDVDQPSPSAQDTQSIA
21245	51613	A	21372	208	423	LYVKPTAPPK*AL*SEGLE*MSP QCLWQGLQQGKAQKKLLEVP NKLTQTIN*PANVLILA*GGSSS YRPPHK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21246	51614	A	21373	1	2002	MPQLPGISLPEGVDPSFLAALPD
						DIRREVLQNQLGIRPPTRTAPST
		İ				NSSAPAVVGNPGVTEVSPEFLA
						ALPPAIQEEVLAQQTAEQQRPE
						LAQNASSDTLMDPVTLIQTLPS
:						DLRRSVLEDMEDSVLAVMPPDI
						AAEAQALRREQEARQRQLMHE
						RLFGHSSTSALSAILRSPAFTSR
						1
						LSGNRGVQYTRLAVQRGGTFQ
1		ļ				MGGSSSH\NRPSGS\NVDTLLRL
						RGRLLLDHEALSCLLVLLFVDE
1						PKLNTSRLHRVLRNLCYHAQT
					ļ	RHWVIRSLLSILQRSSESELCIET
						PKLTTSEEKGKKSSKSCGSSSHE
						NRPLDLLHKMESKSSNQLSW/L
						SSVSMDAALGFR\TNIFQIQRSG
						GRKHTEKHASGGSTVHIHPQA
						APVVCRHVLDTLIQLAKVFPSH
						FTQQRTKETNCESDRERGNKA
						CSPCSSQSSSSGICTDFWDLLVK
1						LDNMNVSRKGKNSVKSVPVSA
						GENKVSEAQANSGSGASSTTTA
						TSTTSTTTTTAASTTPTPPTAPTP
						VTSAPALVAATAISTIVVAASTT
						VTTPTTATTTVSISPTTKGSKSP
						AKVSDGGSSSTDFKMVSSGLTE
						NQLQLSVEVLTSHSCSEEGLED
						AANVLLQLSRRPARMLAMHW
						PSLYNVETEKTLALPNLIALQTS
						PFASGVIAKSSTDRLINQCAPV
	1					WRWR
21247	51615	Α	21374	1	6384	MVSPEPKTGHSIRNWLDELKDL
						PILHAYSNLPSSPAVDLAIHSSK
						EGRMDWTEGQVTGPVVRSAAT
1						SGAGSTTSGVVSGSLGSREINYI
						LRVLGPAACRNPDIFTEVANCC
1						IRIALPAPRGSGTGNGSSRIPRES
1						APEMATAESLVEELSEDAAGG
1						ASPGVELPALGCSELPAAEVSP
					-	TASSKNLETICEYAYCMAMLPE
1						TGLDPYPKRGFLDLTQERIWTD
						IPPSPGNIPTTHPLMVRHADHSS
						LTLGSGSSTT
21248	51616	A	21375	1	1356	2.235357.
21249	51617	A	21376	80	608	
	1	<u>.                                    </u>	1-10.0	l		<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
21250	51618	Α	21377	298	1374	GRGDWPDILNLTQATASWTPR
						WCCLRKPVEAPSGGATEERTLS
						LTSLGLSMPADPCEGGARSCLV
						TESARGGLQNGQSVDVEETLPS
		ł				EPENGALLRSERYQGPRRRACS
		ļ				ETLAESRTAVLQQRRAAQLPG
		1				GPAAAGEQRASPSKARLKEFLQ
		1				QCDREDLVELALPQLAQVVTV
		ł	ļ			YEFLLMKVAESLGITEFLRKKEI
		1				HPDNLGPKHLSRDMDGEQLEG
						ASS\EKREREAAEEGLASVKRP
		1				RREALSND\TT\ESLAANSRGRE
						KPRPLHALAAGTIVSQEEDIVT
		1				VTD/GRGACLRMGPLEGVPLEA
						VESVVTVEPEPSQ*NGVRY*NS
						SDFIQRRSMASNVYFSNVNIAQ
		ŀ				MNLLFIKNNVFL
21251	51619	С	21378	354	530	
21252	51620	Α	21379	1	1272	MVLSEKASGSTLRGCTEERTLS
		ŀ				LTSLGLSMPADPCEGGARSCLV
		ŀ				TESARGGLQFLQQCDREDLVEL
						ALPQLAQVVTVYEFLLMKVAE
		ŀ			]	SLGITEFLRKKEIHPDNLGPKHL
						SRDMDGEQLEGASSEKREREA
						AEEGLASVKRPRREALSNDTTE
						SLAANSRGREKPRPLHALAAES
1						HTTTVSGGNGSVFQAGPQLQA
<b> </b>						LANLEARRGSIGAALSSRDVSG
						LPVYAQSGEPRRLTQAQVAAFP
ł						GENALEHSSAQDTWDSLRSPGF
ţ						CSPLSSGGGAESLPPGGPGHAE
	ļ					AGHLGKVCDFHLNHQQPSPTS
ļ					:	VLPTEVAAPPLEKILSVDSVAV
						DCAYRTVPKPGPQPGPHGSLLT
1						EGCLRSLSGDLNRFPCGMEVHS
		}				GQRELESVVCLSAETMAF*NFP\
		1				MGA\MSYCL\RD\RSRFLFRLPM
		<u> </u>				GLSCPLQVQ
21253	51621	A	21380	90	204	
21254	51622	В	21381	13	423	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21255	51623	A	21382	1513	2405	RRAQYRRWWRRWRRHREPE KEKRRHSRSHSRRRRTLIF*TQS RSRSRE*LHCQSRSR**FKSPRQ RRSHSRERGRSTARHYRSTKER QQRRSRSGTRSPKKSRCPKRKL SLSPSPGKHKKKKKDKERNRY ERERSTSKKKSKDREKDQERKS ESDTDVKVTRVYDEEEQGYDT SQEIEMHADNPAAIQTVVLQRD DLQNGLLSTCRELSRATAELER AWREYDKLEYDVTVTRNQMQ EQLDHLGEVQCWCYRTEKYV MERGPEQDPKRAFLDFVQERIQ GESAVQSESKFIKTVKK
21256	51624	Α	21384	1	180	055 2555
21257	51625	Α	21385	3	115	
21258	51626	Α	21386	3	312	LPTALLRSTMALLHSGRVLPGI AAAFHPGL/AA/AASARASSWW THVEMGPPDPILGVTEAFKRDT NSKKMNLGVGAYRDDNGKPY VLPSVRKFVTVQTISGTGAL
21259	51627	Α	21387	3	395	STMALLHSGRVLPGIAAAFHPC\ LAAAASARASSWWTHVEMGPP DPILGVTEAFKRDTNSKKMNLG VGAYRDDNGKPYVLPSVRKAE AQIAAKNLDLLPLLNVVCSLSS HTVSPAAQGGTRKEDILGFACC
21260	51628	Α	21388	1	303	
21261	51629	Α	21389	2	425	RPE*WKEIATVVKKRNLFALFD MAYQGFASGDGDKDAWALHH FIEQGINVCLCQSYAKNMGLYG EHVGAFTVVCKDADEAKRVES QLKILISPMYSNSPLNGAQIASTI PNTPVGLKVWVMGGLHDHPQ VMSDEKETTLRA

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21262	51630	Α	21390	3	1508	HEAGAGSLCVCPARPLLRPAAA
						HCPSYRSTMAHACTSGRVLPGI
						AAA\FHPGLAAAASARASSWW
						THVEMGPPDPILGVTEAFKRDT
						NSKKMNLGVGAYRDDNGKAL
						PCCPSVRKAEGPRLPQKIWDKE
						YL\PIGG\LAEFCKASAELALGE
ļ						NNRSLERVGRFVTVQTHFWEL
						GALRI\GASFL\QRFFKFSRDVFL
1						PKPTWGETTHPSFRDAG\MQL\
ļ						QGYR\YYDPKTCGFDFTGAVED
ŀ						ISKIPEQSVLL\LHACG/HNNPTG
ŀ						VDPRPEQWKEIATVVKKRNLF
						AFFDMAYQGFASGDGDKDAW
-						AVRHFIEQGINVCLCQSYAKNM
1					]	GLYGERVGAFTMVCKDA\DES
						KRVESQLKILIRPMYSNPPLNG
İ				ļ		ARIAAAILNTPDLRKQWLQEVK
				:		GMADRI\IGHAGLQLVSNLQGR
						EG\STH\NWATHSPTKLGMFCF
						HRG*KLEQVERLIKEFSIYMTK
						DGRISVAGVTSSNVGYLAHAIH
1						QVTKYVSLVRGNGDNLSVLSL
21263	51631	A	21391	1	184	
21264	51632	A	21392	70	253	
21265	51633	Α	21393	1	351	VMEFYCQSCETAMCRECTEGE
1		l				HAEHPTVPLKDVVEQHKASLQ
						VQLDAVNKRWGTLPKPPTGCL
						YLRPGLMASC\QLTNQKASI/VD
						DIHSTFDELQKTLNVRKSVLLM
						ELEVNYGLKHK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21266	51634	Α	21394	208	2546	WLHCMQQQRAGSKTAGPP\CQ
						WSRMAS\EGTNIP\SLVVGQI\DK
						QFLICSICLERYKNPKVLPCLHT
						FCERCLQNYIP\AHSLTLSCPVC
		ŀ				RQTSILPEKGVAALQNNFFITNL
		:				MDVLQ\RTPGSNAEESSILETVT
						AVAAGKP\LSCPNHDGNVMEF
						YCQSCETAMCRECTEGEHAEH
	1					PTVPLKDVVEQHKASLQVQLD
						AVNKRWGTLPKPPTGCLYLRP
	1					GLMASC\QLTNQKASIVDDIHST
	ŀ	1				FDELQKTLNVRKSVLLMELEV
	1					
						NYGLKHKVLQSQLDT\LLQGQE
		1				SIKSCSNFTAQTLNHGTETEVLL
						VKKQMSEKLNELADQDFPLHP
						RENDQLDFIVETEGLKKSIHNL
						GTILTTNAVASETVATGEGLRQ
		1				TIIGQPMSVTITTKDKDGELCKT
						GNAYLTAELSTPDGSVADGEIL
						DNKNGTYEFLYTVQKEGDFTL
						SLSLYDQHIRGSPFKLKVIRSAD
						VSPPTEGVKRRVKSPGSGHVKQ
						KAVKRPASMYSTGKRKENPIED
	ŀ					DLIFRVGTKGRNKGEFTNLQGV
		İ				AASTNGKILIADSNNQCVQIFSN
						DGQFKSRFGIRGRSPGQLQRPT
		İ				GVAVHPSGDIIIADYDNKWVSI
						FSSDGKFKTKIGSGKLMGPKGV
						SVDRNGHIIVVDNKACCVFIFQ
		ļ.				PNGKIVTRFGSRGNGDRQFAGP
		ĺ				HFAAVNSNNEIIITDFHNHSVKV
						FNQEGEFMLKFGSNGEGNGQF
21267	51635	Α	21395	1	591	
21268	51636	Α	21396	3	608	LEGDQRWINTVSQALHLLSTLY
						GPFPNCYGIGRCAKMAYELWR
						NLEEEEDGETKGRRPEIGHIFLL
						DRGKLCPAFLSSFSPRRHGNME
	1	l				GVVAKLKS*S*GCESQL*GAWV
						LGPMESIKAALGHAV*KAGSGR
						TFMVRDLCF\DVDFVTALCSQV
						VYEGLVDDTFRIKCGKCYGPLH
						LGKMDGSSALAPATWGRVCGR
						DPHPE
21269	51637	Α	21397	268	402	
21270	51638	A	21398	139	452	
				<u> </u>		

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
ļ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21271	51639	Α	21399	1	2497	MGPVAAPLGLSLPATDGAHSV
						RGNSVLVPSGLIPKDYRSLKTQ
						YLQSYGPEHLLTFSNLRRAGLL
						TEQAPGDTLTAVESKVSKLVTD
						KAAGKITDAFSSLAKRSNFRAIS
						KKLNLIPRVDGEYDLKVPRDM
		İ				AYVFGGAYVPLSCRIIEQVLER
		ŀ				RSWQGLDEVVRLLNCSDFAFT
		l				DMTKEDKASSESLRLILVVFLG
						GCTFSEISALRFLGREKGYRFIF
		•				LTTAVTNSARLMEAMSELPGK
		ŀ				KDLFIETDLMSPLDRIANVSLLK
İ		l				QHEVDKLYKVENKPALSSNEQ
						LCFLVRPRIKNMRYIASLVNAD
						KLAGRTRKYKVIFSPQKFYACE
						MVLEEEGIYGDVSCDEWAFSLL
	ĺ	ŀ				PLDVDLLSMELPEFFRDYFLEG
						DQRWINTVAQALHLLSTLYGPL
						SQTCYGIGRCAK\MAYEFVEGT
						WEEEEDGRNPRARREGFG\HIF
						LL\DR\DVDFVDSTLLPKLVYEG
						LVDDTFRIKCGSVDFGPEVTSS
						DKSLKVLLNAEDKVFNEIRNEH
		İ				FSNVFGFLSQKARNLQAQYDR
		l			[	RRGMDIKQMKNFVSQELKGLK
						QEHY\LLSLHIG\ACESIMKKKT
						KQDFQE\LIKT\EHALLEGFNIRE
						STSYIEEHIDRQVS\PIESLRLMC
i						LLSITENGLIPKDYRSLKTQYLQ
						SYGPEHLLTFSNLRRAG\LLTEQ
						APGDNLTAVESKVSKL\VTDKA
						AGKITDALS\SLAKRSQFRAISK
21272	51640	A	21400	1	216	NORTH TESTS IN THE STATE OF THE RISK
21273	51641	A	21401	2	270	
21274	51642	A	21402	1	600	SGRALHASWAAGGVGAYSGRL
-12/7	10.72	ľ <b>.</b>	-1.102	-		RSDALEGESFA\LSF\SSASDAEF
						DAVVGYLEDIIMDDEFQLLQRN
						FMDKYYL\EFEDTEENKLIYTP\I
						FNEYISLVEKYIEEQ\LLQR\IPEF
						NMAAFHHNITPSVPVCRHHKD
						EVAW*QYSAMLAPPFTRFSWA
						F*RNVFGTYRGRKKEGRGL\DL
						SSGLV\VTSLCKSSSL\PASQNNL
21275	51642	_	21402	1	207	RH
21275	51643	Α	21403	1	297	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21276	51644	A	21404		1419	MGAPLLSPGWGAGAAGRRWW MLLAPLLPALLLVRPAGALVEG LYCGTRDCYEVLGVSRSAGKA EIARAYRQLARRYHPDRYRPQP GDEGPGRTPQSAEEAFLLVATA YETLKIVFPELKIVKELTTLLMP SRAVWIIWSHKMSGGGAGPEL DIGYVPVLLHMPKRIGSGMDTS RSTTVIPVEIRNKELIALLVKAQ DRQEEETNYKLRPFNGLSYEVG YLVILQDEETRKDYDYMLDHP EEYYSHYYHYYSRRLAPKVDV RVVILVSVCAISVFQFFSWWNS YNKAISYLATVPKYRIQATEIA KQQGLLKKAKAKSKNNKVQR RNS*\EEENIIKNIIKSKIDIKGGY QKPQICDLLLFQIILAPFHLCSYI VWHCRWIYNFNIKGKEYGEEE RLYIIRKSMKMSKSQFDSLEDH QKETFLKRELWIKENYEVYKQ
						EQEEELKKKLANDPRWKRYRR WMKNEGPGRLTFVDD
21277	51645	A	21405	1	1266	WWW.DOI GILLII VDD
21278	51646	A	21406	1	739	
21279	51647	A	21407	1	428	

	SEQ ID NO:	viet	SEQ ID NO:	Nucleotide		Amino acid sequence ( X=Unknown,
	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21280	51648	Α	21408	99	2338	MPEEVHHGEEEVETFAFQAEIA
						QLMSLIINTFYSNKEIFLRELISN
						AS\DALDKIRYESLTDPSKLDSG
						KELKIDIIPNPQEPYPGLWVDTG
						IG\MTKADLINNLGTIAKSGTKA
						FMEALQAGADISMIGQFGDGFY
						S\ANLVAK\KVVVITKHP\DDEQ
1						YAWESSAGGSFTVRADHGEPIA
						\RGT\QVILHLKEDQTEYLEERR
						V\QEVVKKH\SQFIGYPITLYLE
						KEREKEISDDEAEEEKGEKEEE
						DKDDEEKPKIEDVGSDEEDDSG
						KDKKKKTKKIKEKYIDQEELNK
					:	TKPIWTRNPDDIT\QEEYGEFYK
						SLTNDWEDHLAVKHFSVEGQL
						EFRALLFIPRRAPFDLFENKKKK
						NNIKLYVRRVFIMDSCDELVPE\
1						YLNFIPGVVDSEDLPLNIFRKML
						QQNK/ILWKVIRKNIVKKCLELF
						SELAEDKENYKKFYEAFSKNLK
						LGIHEDSTN\RRRLSELLRYHTS
						QSG\DEMTSLSEYVS\RMKETQ
						MSIYYITGESK\EQVANSAFVER
						VRKRGFEVVYMT\EPI\DEYCVQ
						QLKEFDGKKLV\SVTKEGLELP
						EDEEEKKKMEESKAKFENLCK
İ						LMKEILDKKVEK\VTIS\NRL\VS
						SPC\CIGHKHLGAGQALIERDHE
						KPRALRDHPPQGGIMMGKKAP
						*ESPPDHPHFVETRAGRKADAR
						QVMIRAVKD/LWWVLLV*NPP
						LLSSGF\SLEDPQT/HTPNRIYRI
	51649	Α	21409	1	426	
	51650	A	21410	1	1305	
21283	51651	A :	21411	446	3180	VAVRLGSASQFPSGARPQLPRP
						PSWSSSIHSPISTTGLSESTDSDS
		'				KSGLRGFSAYNPSSRIPKSTTSS
						NSSDGAVTASNQDTGIPWVLV
						TGPTPPVNSPVESLQGSAFSSGF
						STPRGLVKGPTLPPDSNLTSVSD
						YTWIFVRGPPIPNDLSQVSIQAP
						PTPSNTPECQYKGLPHPQCPASP
						GVWYQDHLYPITFPQCRHKHL
						LHPITSPQCQGRRLPPPQCLTLR
						GVQYKNLLHPVILSSCRHKDIL
<u> </u>		<u> </u>				NLITLAE

SEQ ID	SEQ ID NO:		SEQ ID NO:	Nucleotide		Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21284	51652	A	21412	24	667	MLPWLNCITTGVPWTPTSKRW
İ						LRNSKVCDCCDKTPSNAFSLLS
						VPPTTTSPASLAWWSGCARLLD
						LGSSSLMMSPTMASPACRPWL
						GQRWRLISGSWAWAIVPVT*VP
1						VPEPSWKNRAG*PGCSSYESPH
						MRRPTRPSASCLEWAPRASLA
						MHLMATSFCVLLSSARITSEKA
						PLSGEGIHKVKNWNPSFPPASPP
						FPMGSVTTAGPRTWMTLP
21285	51653	Α	21413	1	304	RAQVLPARCYPGSTVSPLGFRG
						LPLPRGGSEIPRCATAATRPHR
						MPFLFYLFLQQQHRPHHWHGG
						AAVPGFWTSAHPA\CDVTYHGF
						PSLQALAGPPWQCI
21286	51654	Α	21414	1	7629	
21287	51655	A	21415	180	423	
21288	51656	Α	21416	17	753	QAHTRRAGGRAQVLPARCYPG
						STVSPLGFRGLPLPRGGSEIPRC
		Ì				ATAATRPHRMPFLFYLFLQQQH
						RPHHWHGGAAVPGFWTSAHPA
						\CDVTYHGFPSLQALAGPEVEA
						HLRKLGLGYRARYVSASARAIL
						EEQGGL/RLAAAATRVLI*GGP
				•		QGPLHPAWSGHQGG*LHLPDG
						PRQAPGCARGCPYVAHCPT*LQ
				:		LAPYHVPGEGTEPPDQQGTGKL
						FPEPVGTLCWLGPSALSGHHHF
						YDLSRTP
21289	51657	Α	21417	17	1087	QAHTRRAGGRAQVLPARCYPG
		ŀ				STVSPLGFRGLPLPRGGSEIPRC
						ATAATRPHRMPFLFYLFLQQQH
İ						RPHHWHGGAAVPGFWTSAHPA
		·				\CDVTYHGFPSLQALAGPEVEA
						HLRKLGLGYRARYVSASARAIL
						EEQGGL/RLAAAATRVLI*GGP
						QGPLHPAWSGHQGG*LHLPDG
						PRQAPGCARGCPYVAHCPT*LQ
						LAPYHVPGEGTEPPDQQGTGKL
						FPEPVGTLCWLGPSGWIRCLLK
						NYRLLPLDLEGLLGNAFDGHQ
						LLRPLIFCQDHLREGPPIGRGDS
						QGEELEPQLPSSLSSIPYGFCDH
						CWTKDVDDPPLVTHPSPGSRD
						GHMTQAWPVKVVSPLATVIGH
						VMQASLLAL
<u> </u>	L		L	<u> </u>	1	

NO: of peptide hod in USSN location of first codon for last amino acid *=S	nino acid sequence ( X=Unknown, Stop codon, /=possible nucleotide letion, \=possible nucleotide insertion)
EG RK HR RR AR AS WF TQ TPJ YH VR RITI VT KL GG CIL QA PT SL'	QKNTAVRAHRLWGREKISRK GAGPTPQESRRIGARSGALMT KGRGMQEVEELSETGKVVKQ RVGEALRVVVLVWAGSLG/D RGLVLGRRGYYGAVPAVEMP RALLPRRMGHRTLASTPALW SIPCPRSELRLDLVLPSGQSFR (REQSPAHWSGVLADQVWTL QTEEQLHCTVYRGDKSQASRP PDELEAVRKYFQLDVTLAQL HHWGSVDSHFQEVAQKFQG RLLRQDPIECLFSFICSSNNNIA ITGMVERLCQAFGPRLIQLDD TYHGFPSLQALAGPEVEAHLR LGLGYRARYVSASARAILEEQ GLAWLQQLRESSYEEAHKAL ILPGVGTKVADCICLMALDKP AVPVDVHMWHIAQRDYSWH ITSQAKGPSPQTNKELGNFFR LWGPYAGWAQAGLLGNAFD HQLLRPLIFCQDHLREGPPIGR DSQGEELEPQLPSSLSSIPYGF DHCWTKDVDDPPLVTHPSPG RDGHMTQAWPVKVVSPLATV GHVMQASLLAL

2717

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21291	51659	Α	21419	837	2785	WTRMLFAVSCNDGECQNSNVI
						LFCDMCNLAVHQECYGVPYIPE
						GQWLCRRCLQSPSRAVDCALC
						PNKGGAFKQTDDGRWAHVVC
						ALWIPEVCFANTVFLEPIDSIEHI
						PPARWKLTCYICKQRGSGACIQ
•						CHKANCYTAFHVTCAQQAGLY
					:	MKMEPVRETGANGTSFSVRKT
						AYCDIHTPPGSARRLPALSHSE
						GEEDEDEEEDEGKGWSSEKVK
						KAKAKSRIKMKKARKILAEKR
i						AAAPVVSVPCIPPHRLSKITNRL
						TIQRKSQFMQRLHSYWTLKRQ
						SRNGVPLLRRLQTHLQSQRNCD
						QVGRDSEDKNWALKEQLKSW
						QRLRHDLERARLLVELIRKREK
						LKRETIKVQQIAMEMQLTPFLIL
						LRKTLEQLQEKDTGNIFSEPVPL
					:	SEVPDYLDHIKKPMDFFTMKQ
1		ŀ				NLEAYRYLNFDDFEEDFNLIVS
						NCLKYNAKDTIFYRAAVRLRE
						QGGAVVRQARRQAEKMGIDFE
[						TGMHIPHSLAGDEATHHTEDA
<u> </u>						AEEERLVLLENQKHLPVEEQLK
						LLLERLDEVNASKQSVGRSRRA
						KMIKKEMTALRRKLAHQRETG
						RDGPERHGPSSRGSLTPHPAAC
						DKDGQTDSAAEESSSQETSKGL
						GPNMSSTPAHEVG\GEPQFCSP
	]					KRTRRQ/HGPPKRPGRPPKNRES
						QMTPSHGGSPVGPP/RAPHHEFP
	-					ASAQAG*EPQFCSPKRTRRQLD
21292	51660	Α	21420	1	2625	MENKITGYTTVDISQWHRKEHF
						EAFQSVAQCTYNQTVQLDITAF
						LKTVKKNKHKFYPAFIHILARL
ŀ						MNAHPEFRMAMKDGELVIWD
						SVHPCYTVFHEQTETFSSLWSE
ł						YHDDFRQFLHIYSQDVACYGE
						NLAYFPKGFIENMFFVSANPWV
						SFTSFDLNVANMDNFFAPVFTM
						GKYYTQGDKVLMPLAIQRDSE
						DKNWALKEQLKSWQRLRHDL
						ERARLLVELIRKREKLKRETIKV
1						QQIAMEMQLTPFLILLRK

SEQ ID	1 `		SEQ ID NO:	l	ľ	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				Sequence		
21293	51661	Α	21421	164	1401	PPKGRGMQEVEELSETGKVVK
						QHRVGEALRVVVLVWAGSLG/
						DRRGLVLGRRGYYGAVPAVE
						MPARALLPRRMGHRTLASTPA
						LWASIPCPRSELRLDLVLPSGQS
		İ				FRWWREQSPAHWSGVLADQV
						WTLTQTEEQLHCTVYRGDKSQ
						ASRPTPDELEAVRKYFQLDVTL
						AQLYHHWGSVDSHFQEVAQKF
						QGVRLLRQDPIECLFSFICSSNN
						NIARITGMVERLCQAFGPRLIQL
						DDVTYHGFPSLQALAGPEVEA
						HLRKLGLGYRARYVSASARAIL
						EEQGGLAWLQQLRESSYEEAH
						KALCILPGVGTKVADCICLMAL
						DKPQAVPVDVHMWHIAQRDY
						SWHPTTSQAKGPSPQTNKELGN
						FFRSLWEPYAGWAQATPPSLQ
						VLFSADLRQSRHAQEPPAKRRK
21294	51662	Α	21422	751	2107	SSEHPRELHRIPISWAMSELKDC
						PLQFHD\FKSVDHLKVCPRYTA
						VLARS\EDDGIGIEELDTLQLEL
İ						ETLLSSASRR\LRVLEAETQILT
						D\WQDK\KGDRRFLKLG\RDH\E
						FGAPPKHGKPKK\QKLE\GKAG
İ						HGPGP\CPGRPKSKNLQPK\IQE
						YEFTDDPIDVPRIPKNDAPNRF
						WASVEPYCADITSEEVRTLEEL
ł		1	ļ			LKPPEDEAEHYKIPPLGKHYSQ
						RWAQEDLLEEQKDGARAAAV
	1					ADKKKGLMGPLTELDTKDVDA
		İ				LLKKSEAQHEQPEDGCPFGALT
						QRLLQALVEENIISPMEDSPIPD
						MSGKESGADGASTSPRNQNKP
						FSVPHTKSLESRIKEELIA\QALL
						*AGDLLAKDSEDEVLAELRKR
		1		ļ		QG\EL\KALSAHNRTKKA\DLLR
						LAKEEVSRQE\LRQR\VRRAERE
		1				GMDAFRKIHGLPGRRKRNSPPK
						K\EKDQAWKNS*KEREEHP
21295	51663	Α	21423	107	331	

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
		<u> </u>				
21296	51664	Α	21424	387	1071	NNPEWRTLIYVDKENGEPGTR
ŀ						VVAKDGLKLG\SGPSIKALDG\R
ŀ						SQVS\TPRFGKTFDAPTQPYLKL
						PRKALG\TVNR\TTEKSVKDPGD
						PSNKKQPKLFRPKKDGPEEGLL
ŀ						KAKKPLFPGLQD*CPIPEIRKIPF
1						PFNPS*TLRSF*PCPEEHPDCGTF
1						PL*VGV\PLMIPLTREREL*KSW
						FQLGPPFHLWKDGPLPPWGIPIC
						LQFSFQGISVRTWGV\DLPPVCC
						DIDI
21297	51665	Α	21425	480	1270	RSVKKVGAQRPGSGGGWGGN
						EGWVPQATAPGWAYTHVLFLY
1						LFVTTPISAALRADGFGDNPFY
						HCLVAEILPAPGKLLGKSTWPL
						EVHLSSRPQRSPHRLPEALLPFS
		l				LSQG/HCVVGYGIYYFIYSTWK
						GRTIYLEDIYVMPEY/RGYWAE
						AEAGVGAKHPTDPTVIPCPQSP
						AGVIF*SPSAQTIAIFCLFLSTTG
	:					QGIGSKIIKKVAEVALDKGCSQ
						FRLAVLDWNQRAMDLYKALG
						AQDLTEAEGWHFFCFQGEATR KLAGK
21298	51666	A	21426	182	<b>7</b> 97	LPFARPWDFGPRQRGSGWLPC
21298	131000	A	21420	102	191	GSREAKEGDCGDIL\RLIRVKTA
						GPELAEFEKLFGSGERSVEEAL
						ESRLAFGDNPFYHLFW*AEILP
		İ				APGESYWG/HCVVGYGIYYFIY
1						STWKGRTIYLEDIYVIAGNIGV
						QGIGSQK*SKKV\AEVALGIRGC
						SPNFRLARPLD\WNPEGPWDLY
						KAPRSPKI*RKLEGWHFFCFQG
						EATRKLAGK
21299	51667	Α	21427	224	490	QAGFISARNITWR*ANRRSTWR
12,233	31007	ľ .	- · · · ·			SYRRKSGRTANCPPISPPACSRC
						SAPIFIPKNGWWRAMVAPLKW
						PSTGAFNRLFVPLLTWWRSPDR
21300	51668	A	21428	301	773	HGPGTQQRPRSIWRMPQ*ARPG
1300	0001	^\	21420	301	113	GRSLAVGGGPRPRAAGGFP*VA
			[			SRARRCQR\AHGCHCAFSPLSP
			1			ALLSQDAGLCQSRPLLLGQRRC
	1					LSQEVWSGLGNVHFCQPPEDA
						ATTPSCPEALTAQGQQASVCDR
						VSHRFSTWSMVRCSDRTEGQR
1						LQTLGD
21301	51669	В	21429	1	1380	r X I r O D
21301	21003	1 <sub>D</sub>	21727	I	1:300	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21302	51670	A	21430		1660	AGVLASHPNSTDVHIINLSLTFH GQELLSDTKLELNSGRRYGLIG LNGIGKSMLLSAIGKREVPIPEH IDIYHLTREMPPSDKTPLHCVM EVDTERAMLEKEAERLAHEDA ECEKLMELYERLEELDADKAE MRASRILHGLGFTPAMQRKKL KDFSGGWRMRVALARALFIRPF MLLLDEPTNHLDLDACVWLEE ELKTFKRILVLVSHSQDFLNGV CTNIIHMHNKKLKYYTGNYDQ YVKTRLELEENQMKRFHWEQD QIAHMKNYIARFGHGSAKLAR QAQSKEKTLQKMMASGLTERV VSDKTLSFYFPPCGKIPPPVIMV QNVSFKYTKDGPCIYNNLEFGI DLDTRVALVGPNGAGKSTLLK LLTGELLPTDGMIRKHSHVKIG RYHQHLQEQLDLDLSPLEYMM KCYPEIKEKEEMRKI\IGRYGLT G\KQQVSPIRNLSDGQKCRVCL AWLAWQNPHMLFLDEPTNHL DIETIDALADAINEFEGGMMLV SHDFRLIQQVAQEIWVCEKQTI TKWPG\DIL\AYKEHLKSKLVD\

2721

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	location of first	codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
21303	51671	Α	21431	78	2022	NNSNLPLWVIIMPSDLAKKKAA
			-			KKKEAAKARQRPRKGHEENGD
	1					VVTEPQVAEKNEANGRE/TTRE
		ŀ				VDLLTKELEDFEMKKAAARAV
						TGVLASHPNSTDVHIINLSLTFH
		ŀ		:		GQELLSDTKLELNSGRRY\GLIG
		ŀ				LNGIGIGKSMLLFAIGKREVPIP
				:		EHIDIYHLTREMPPSDKTPLHCV
		ł				MEVDTERAMLENEAERLAHED
		Ì				AECEKLMELYERLEELDAD*AE
		ŀ				MRASRI\LHGLGFTPAMQRKKL
						KDFSGGWRMRVALARALFIRPF
						MLLLDEPTNHLDLDACVWLEE
		į		1		ELKTFKRILVLVSHSQDFLNGV
				Ì		CTNIIHMHNKKLKYYTGNYDQ
						YVKTRLELEENQMKRFHWEQD
						QIAHMKNYIARFGHGSAKLAR
				-		QAQSKEKTLQKMMASGLTERV
						VSDKTLSFYFPPCGKIPPPVIMV
						QNVSFKYTKDGPCIYNNLEFGI
						DLDTRVALVGPNGAGKSTLLK
						LLTGELLPTDGMIRKHSHVKIG
						RYHQHLQEQLDLDLSPLEYMM
		1				KCYPEIKEKEEMRKIIGRYGLT
						GKQQVSPIRNLSDGQKCRVCLA
						WLAWQNPHMLFLDEPTNHLDI
						ETIDALADAINEFEGGMMLVSH
					}	DFRLIQQVAQEIWVCEKQTITK
						WPGDILAYKEHLKSKLVDEEPQ
						LTKRTHNVCTLTLASLPRP
21304	51672	A	21432	313	599	
21305	51673	Α	21433	737	1006	DNPVVFVFGSVYMLDYVY*FS
						YVEPPLHPWDEAHLIMVDKLF
						DVLLDSVCQYFIEDFCINVNQG
		L				YWSKILFFCCISARLWYQDDAG
21306	51674	Α	21434	1108	1260	GVNF\NLSCFLLWAFSAIHFPLH
						TALNVSQRFWYVVSLFSLVSK
						NIFISAF
21307	51675	Α	21435	804	917	
21308	51676	A	21436	158	298	
21309	51677	A	21437	1	1068	
21310	51678	A	21438	1	1188	
21311	51679	A	21439	122	352	ICCVICECEVI ELI*CICI EWYOY
21312	51680	Α	21440	647	662	ICGVISEGSVLFH*SISLFWYQY
						HAVLVT/DSLVV*FEVT*HDASS
21212	51601	D	21441	53	692	FVLLA*DRRPEGSSPKSP
21313	51681	В	21441	1	925	
21314 21315	51682 51683	B A	21442	606	900	
21313	121003	IA.	Z1443	1000	200	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	,	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21316	51684	A	21444	33	852	CWKFWARAN*AGGITLPDFKL
						YYKATVTKTAWKFHFTMFGV
						DLKETCTAPSLWKDLALNTVE
						LVCKLCVRQVEGARQIFQLNCT
						VSE
21317	51685	A	21445	3	795	
21318	51686	Α	21446	475	1023	CWFSAGDGGLPGPDYGQDFAA
						HPAGAGALYKCRDLINDNILRT
				į		VSCQSHDDGDIRGMAFARQ*Q
						RPVQIDHNMARRG**AATFQPI
		l				NKTFSGIHWPHGMRTGGTNPN
						FENIECTNHIHTLGLPFSEVERN
						DCDGFNFQLHIFRQTRDLHTAP
						GRIGRLMLIEELLIDFVDGGKII
				•		QIGDKHGH
21319	51687	Α	21447	1637	2020	PSSEEFTGGPARGFPERARAQL
						TGRTPGQRTQLDNDLLTKSTP
				:		WRGVMGPTLLPGLLAGPFQPL
		ł				GHTGPFYPQGFRWQKTHWVRL
						AQAHWQF*LGQFRGPKTADGG
						GSKGPGLSASHGAKHHVITYV
21320	51688	Α	21448	1051	1185	
21321	51689	A	21449	1290	2025	
21322	51690	В	21450	1	1410	
21323	51691	Α	21451	3269	3678	SLSIRLATAPCWLRYFASSRRFQ
		İ				PTRSAGSVTDSSVTASASSPG*R
						*NASP\CRERSHRSSRGQARRSL
		l				HLWDRSDR*TLPCDHGRCCHA
						NDQQVGFAGNYSTAGQT*RWT
		ŀ				GFTAGHAGRDINKSADKLPAT
		ļ				AGKR
21324	51692	В	21452	2655	2825	
21325	51693	A	21453	[1	896	MGDFNTPLSTLDRSMRQKVNK
						DTQELNSALHQADLIDIYRTLH
						PKSTEYTFFSAPHHTYSKIDHIL
				İ		GSKALLSKCKRTEIITNYLSDHS
						AIKLELRIKNLTQNHSTTWKLN
				ļ		NLLLSDYWVHNEMKAEIKMFF
		:				ETNENKDTTYQNLWDAFKAVC
						RGKFIALNAHKRKQERSKIDTL
						TSQLKELEKQEQTHSKASRRQE
			1			ITKIRAELKEIETQKTLQKINESR
1						SWFFERINKIDRPLARLIKKKRE
		1				KNQIDTIKNDKGDITTDPTE\YK
1						LPSENTTNTSTQIN*KI*KKWIN
		<u> </u>				SSTHTLSQD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21326	51694	A	21454	23	559	TPGLRQSSLSPSAFASQNAGVS
						GIGGFLVSLTSRMKPWTLAMFR
						CVWSFFLLVGSWSCWLQE*SC
						RPSQ*VSQFL/NSSASGVVCCSC
						LELFVPPGGFVVSLASRVKLQT
			ŀ			FTVSVTAHKGGVDPKSEQQQD
						LVQKAKEHSFCSVEADLSGLPL
	! !					LAGAACFYSLMEPHPHPADWSI
						LQRAD
21327	51695	A	21455	26	323	
21328	51696	Α	21456	221	734	
21329	51697	A	21457	293	2103	PCHSALSSPLPHRDCIALCFLNS
ļ						GTSFVAGFAIFSILG\FMF\QEQG
						VPISEVAESGPGLAFIAYPRAVV
						MLPFSPLWACCFFFMVVLLGL
						DSQFVCVESLVTALVDMYPHV
ľ						FRKKNRREVLILGVSVISFLVGL
						IMLTEGGMYVFQLFDYYAASG
						MCLLFVAIFESLCVAWVYGAK
						RFYDNIEDMIGYRPWPLIKYCW
						LFLTPAVCTATFLFSLIKYTPLT
						YNKKYTYPWWGDALGWLLAL
						SSMVCIPAWSLYRLGTLKGPFR
						EVIIPVSIRMTKVAGTAGLSGN
						VGPRCDLSRIEELPDGGTRCGM
						DSRVSGTTSNGETKPVYPVME
						KKEEDGTLERGHWNNKMEFVL
		1				SVAGEIIGLGNVWRFPYLCYKN
						GGVVGFYNSIKNIPILQRVILFV
						RIHNMLNKGQGVLAKCDAAK
	:	1				KIEEKEADAGSASVPGDVMRT
						KKLSFEGAWHPCLLVGLKLSD
						ALELRGGFISARIAGPRAAESDS
						VGLERELRVCIANKLPGEADAA
						GPGTTFWKHCSMPITDGADAV
						EGRNTQEAFFRQLNPEAAFLHF
						MEMVDTNALHVHGIHTRKNHP
						GQKEAAHSQHGTFPIRWTLNLL
						KRTFQKGNRAEPNSPAM
21330	51698	Α	21458	1	419	MKLRTLAVSATALKVARLEFV
						PFDVRMCSEFLSSGVKLQTFAV
						SVTALNALRLELFVPPGGLMVS
						LASGVKLQIFTVSVTAHKSSVD
						PKTLG/CVDGTGRRGAGGGAH
						RGGLGRTGAHGGGGRLRHGGL
}						QVSSPAPRKGS

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	in USSN	location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
21331	51699	A	21459	24	1639	GGMDSRVSGTTSNGETKPVYP
						VMEKKEEDGTLERGHWNNKM
						EFVLSVAGEIIGLGNVWRFPYL
						CYKNGGGAFFIPYLVFLFTCGIP
						VFLLETALGQYTSQGGVTAWR
						KICPIFEGIGYASQMIVILLNVY
						YIIVLAWALFYLFSSFTIDLPWG
						GCYHEWNTGNWVDVVE/HGG*
						GEWCGNGSNTPHVCR/VTGWT
						W*SRWLKGMVWEWIQHTPCV
						QGTGWTW*STWLKG/PGNRVD
						VVEHVAEGNGVGMDPTHPTCA
						GNRVDVVEQVAEGNEHCMEF
						QKTNGSLNGTSENATSPVIEFW
						DSPPGPQFVCVECLVTASIDMF
		İ				PRQLRKSGRRELLILTIAVMCY
						LIGLFLVTEGGMYIFQLFDYYA
						SSGICLLFLSLFEVVCISWVYGA
						DRFYDNIEDMIGYRPWPLVKIS
						WLFLTPGLCLATFLFSLSKYTPL
						KYNNVYVYPPWGY\SIGWFLA
		l				LSSMVCVPLFVVITLLKTRGPFR
						KVGGVGESDLLKEGTERGWT
						WGRTHNWGQGADTNSECGAA
21222	61700	<u> </u>	21460	1250	4707	LQSHRILTRRTLKTSPVAFQKCS
21332	51700	A	21460	1358	4787	RRCYCGIWSRYLLKRLWKWIS
						GEGWAEKDSARRRGICTFWLE FE/VTCLSWNSLF/YAHINVDISG
						MKVPWLYVGMCFSSFCWHIED
						HWSYSINYLHWGEPKTWYGVP
						SHAAEQLEEVMRELAPELFESQ
						PDLLHQLVTIMNPNVLMEHGV
						PVYRTNQCAGEFVVTFPRAYHS
						GFNQGYNFAEAVNFCTADWLP
						IGROCVNHYRRLRRHCVFSHEE
İ						LIFKMAADPECLDVGLAAMVC
						KELTLMTEEETRLRESVV
21333	51701	Α	21461	1	5773	MEEKPLKVKGKDSSEKKRKRK
						LEKVEQLFGEGKQKSKELKKM
						DKPRKKKLKLGADKSKELNKL
						AKKLAKEEERKKKKEKAAAAK
				ļ		VELVKESTEKKREKKVLDIPSK
						YDWSGAEESDDENAVCAAQN
						CQRPCKDKVDWVQCDGGCDE
}						WFHQVCVGVSPEMAENEDYICI
1						NCAKKQGPVSPGSHKAVITVV
1						VKAVSYLKFEEESTSELMWLL
						AAFSSFWVVGIKALDFVWPLA
						REEKEIGDNKDLERNKIGVNM

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21334	51702	Α	21462	85	1333	STSRALSGAQRFGRLIDHGARV
	3.702	ľ.	127.02			PALLLLLPL\VHVSATTPEPR\E
						LDDEDFRCVCNFSEPQPDWSEA
						FQCVSA\VEVEIHAG\GLNL\EPF
						LK\RVDADADPRQYA\DTVKGF
		i				PR/VRRASQVGAAQVPA\QLLV
						GALR\VLA\YSRFKELTLEDLKI
						TG\TMPPLPLEATGLALS\SLRLR
						NVSW\ATGRSWL\AEL\QQWLK
						PG\LKVLSIAQAHSPAFSCEQVR
						CFPGPSPA*DLSDNPGTGANAE
						LMAALCPHKFPA\IQNL\ALRNT
					ľ	GNGRRPQGVCAALAAAGVQPH
						SLDVSHNSLRANRNPS\AP\RCM
				İ		\WSNA\LNSPQICRFAGAGTGVP
						KGPAPAKL\RVLRSSACNRL\NR
						A\PQPWTSLPE\VDNLGQLDGNP
						FPGSPGTGPSPTEGSMN\SGVVP
						SLCNVRTCRVGVSGTLVLLQG
						ARGFA
21335	51703	Α	21463	1640	5804	TITGSSPLAGGHHSTCRGRMDM
21333	31703	` `	21.03	1010	3001	VENADSLQAQERKDILMKYDK
			<u> </u>			GHRAGLPEDKGPEPVGINSSIDR
						FGILHETELPPVTAREAKKIRRE
						MTRTSKWMEMLGEWETYKHS
						SKLIDRVYKGIPMNIRGPVWSV
						LLNIQEIKLKNPGRYQIMKERG
į.						KRSSEHIHHIDLDVRTTLRNHV
						FFRDRYGAKQRELFYILLAYSE
:						YNPEVGYCRDLSHITALFLLYL
						PEEDAFWALVQLLASERHSLPG
i						FHSPNGGTVQGLQD
21336	51704	A	21464	1	960	· · · · · · · · · · · · · · · · · · ·
21337	51705	Α	21465	1	1317	
21338	51706	Α	21466	1	3393	
21339	51707	A	21467	2745	6141	QGSKGTCHPQAQQPWDEGVW
						QEAPSQSEPWGQSQEPPTMPQR
						LPHARQHTPLPLGSADYRRVVS
						VRPQGPHRDPKDSRDAAKREQ
						GSLAPRPVPASRGGKTLCKGYR
						QAPPGPPAQFQRPICSASPPWAS
	1					RFSTPCPGGAVREDTYPVGTQG
						VPSLALAQGGPQGSWRFLEWK
						SMPRLPTDLDIGGPWFPHYDFE
						RSCWVRAISQEDQLATCWQAE
						HCGEVHNKDMSWPEEMSFTAN
						SSKIDRQKVPTEKGATGL
21340	51708	Α	21468	65	320	
				3	505	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21342	51710	A	21470	2	1756	CRCHSELSGSSL*/CCTRGIGGAI
						GAVGFRSQAASTLAGQPDGNG
						VCVHAGNSRIFVPLPVGLEVKG
į.						FSHSAFTFGIESHISQSNINGTLV
						PPSALISILQKGLQYVEAEISINK
						DGTVFDSRPIESLSLIVAVIPDV
						VQMRQQAFGEKLTQQQASAA
						ATEASAMAKAATMTPAAISQQ
		İ				NPPKNREATVNGEENGAHEINN
						HSKPMEIDGDVEIPPNKATVLR
		ŀ				GHESEVFICAWNPVSDLLASGS
						GDSTARIWNLNENSNGGSTQL
						VLRHCIREGGHDVPSNKDVTSL
		:				DWNSDGTLLAMGSYDGFARIW
						TENAPALDVDWQNNMTFASCS
						TDMCIHVCRLGCDHPVKTFQG
						HTFCTCIESICFWGGLRKLTTM
						TEGKRLRPKTFCSDGGALLPPA
}						GRRPHLLTGPDIFFKRLSALFQN
						EVNAIKWDPSGMLLASCSDDM
						TLKIWSMKQDACVHDLQAHSK
:		1				EIYTIKWSPTGPATSNPNSSIML
						ASASFDSTVRLWDVEQGVCTH
						TLMKHQEPVYSVAFSPDGKYL
						ASGSFDKYVHIWNTQSGSLVHS
]						YQGTGGIFEVCWNARGDKVGA
						SASDGS\VCILWQLSGKQ
21343	51711	Α	21471	1	1647	
21344	51712	В	21472	4	267	
21345	51713	Α	21473	1	1824	
21346	51714	A	21474	33	543	TFEGANNFVGDPIVLPCNPPKG
	1					LPPLHIYWMNIELEHIEQDERV
	1					YMSQKGDLYFANVGRKRDSRN
•	1					D\YCCFA\AFPRIKGLLYRKMPM
ļ						/RNLTVNSSNSIKAKENPKLLLA
						SPLRSGSESSITILKGEILLLECF
		]				AEGLPTPQVDWNKIGGDLPKG
						REAKENYGKTLKIENVY
21347	51715	A	21475	197	516	TYLAAGRTPTLDPWNQERCTM
						PSGSIWILETTHSSFGWVTVAH
			1			LSSDPKGCQF*VGSRIGEGSATD
		-				PAFCASCSATWTIRPSRSNGA*G
			<u> </u>			VNG**VCWLEPLAGPHR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
21348	51716	Α	21476	1	1766	MNVALTHIDTHHAEFEMALQA
						ALVWNTLRTAGGKKSVHLASA
						LLVHEIIIDGANLTISNVTLEDQ
						GIYCCSAHTALDSAADITQVTV
1						LDVPDPPENLHLSERQNRSVRL
						TWEAGADHNSNISEYIVEFEGN
						KEEPGRWEELTRVQGKKTTVIL
į						PLAPFVRYQFRVIAVNEVGRSQ
						PSQPSDHHETPPAAPDRNPQNIR
						VQASQPKEMIIKWEPLKSMEQN
						GP\GLEYRVTWKPQGAPVEWE
						EETVTNHTLRVMTPAVYAPYD
						VKVQAINQLGSGPDPQSVTLYS
						GEDLPEQPTFLKVIKVDKDTAT
						LSWGLPKKLNGNLTGYLLQYQ
		1				IINDTYEIGELNDINITTPSKPSW
						HLSNLNATTKYKFYLRACTSQ
						GCGKPITEESSTLGEGSVSLEKC
						ENKLIRNIGTKKQGTAIRIPENV
						EAALELRKQQKLEECGGFRRRE
						NANEQQGVTEEFKQQNEKLRF
						LLLKIVILAVAREMTWNGEGR
						DLSEPSLTVVFADLSSPAYSLHT
ĺ						TMAQEPGPGLFLSLSTIPWPISSI
						PMIHASVQVSWIPLLRNSLMVH
1		ĺ		}		IHSSLEQRNLKVHISKEQKKLNS
						VDSDLTDNPPTHTVN
21349	51717	Α	21477	1	1440	
21350	51718	Α	21478	266	3974	RAMEPLLLGRGLIVYLMFLLLK
						FSKAIEIPSSVQQVPTIIKQSKVQ
						VAFPFDEYFQIECEAKGNPEPTF
						SWTKDGNPFYFTDHRIIPSNNS
						GTFRIPNEGHISHFQGKYRCFAS
						NKLGIAMSEEIEFIVPSVPKFPK
						EKIDPLEVEEGDPIVLPCNPPKG
						LPPLHIYWMNIELEHIEQDERV
1						YMSQKGDLYFANVEEKDSRND
						YCCFAAFPRLRTIVQKMPMKLT
						VNSLKHANDSSSSTEIGSKANSI
			<u></u>			KQRKPKL

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21351	51719	A	21479		2421	QAGAMASGSVAECLQQETTCP VCLQYFAEPMMLDCGHNICCA CLARCWGTAETNVSCPQCRETF PQRHMRPNRHLANVTQLVKQL RTERPSGPGGEMGVCEKHREPL KLYCEEDQMPICVVCDRSREHR GHSVLPLEEAVEGFKEQIQNQL DHLKRVTRLKFTDFLAQGEQA RAELLSLTHMEREKIVWEFEQL YHSLKEHEYRLLARLEELDLAI YNSINGAITQFSCNISHLSSLIAQ LEEKHQQPTRELLQDIGDTLSK AERI\RIPEPWITPPDLQEKIHIFA PKCLFLTESLKQFTEKMQSDME KIQELREAQLYSGGSIVGGHEP GEPRGIKAGYGTCNCFPEEEKC FCEPEDIQDPLCDELCRTVIAAA VLFSFVVSVLLSAFCIHCYHKF AHKPPISSAEMTFRRPAQAFPVS YSSSGARRPSLDSMENQVSVDA FKILEDPKWEFPRKNLVLGKTL GEGEFGKVVKATAFHLKGRAG YTTVAVKMLKENASPSELRDL LSEFNVLKQVNHPHVIKLYGAC SQDGPLLLIVEYAKYGSLRGFL
						RESRKVGPGYLGSGGSRNSSSL DHPDERALTMGDLISFAWQISQ GMQYLAEMKLVHRDLAARNIL VAEGRKMKISDFGLSRDVYEE DPYVKRSQGRIPVKWMAIESLF DHIYTTQSDVWSFGVLLWEIVT LGGNPYPGIPPERLFNLLKTGH

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			:	sequence		
21352	51720	Α	21480	79	2253	PSATLSCPTSVAASLGEPAVFPL
						SLRGHRRRHRPGHSKPSVRGRV
						WDTGPGGAALRPPGASQEQRT
						GESAGELGSVQSSAPGPMPVRP
					:	RRPGAMASGSVAECLQQETTC
						PVCLQYFAEPMMLDCGHNICC
						ACLARC\WGTEET\NVS\CPQCR
			İ			E\TFPQRHMRPNRHLANVTQLV
						KQLRTERP\SGPGGEMG\VCE\K
						HREPLKLYCEEDQMPICVVCDR
						SREHRGHSVLPLEEAVEGFKEQ
						IQNQLDHLKRVKDLKKRRRAQ
						GEQARAELLSL\TQ\MEREKIVW
1						E\FEQLYHSLKEHEYRLLARLEE
1						LDLAIYNSINGAITQFSCNISHLS
		l				SLIAQLEEKQQQPT\RELLQDIG
						DTLSRAERIRIPEPWITPPDLQE
						KIHIF\AQK\CLFLTESLKQFTEK
					:	MQSDMEKIQELREAQLYSVDV
						TLDPDTAYPSLILSDNLRQVRY
						SYLQQDLPDNPERFNL\FPCVLG
						SP\CFIAGETIIWEGRGWGDKAK
						WTIG\VC*RLSCQEKGGVTSAP
						QNGFW\AVSLWYGK\EYW\ALT
						SPMTALPLRTPFQ\RVGIFL\DYD
1						A\GEVSF\YN\VTERC/HTPFTFSS
						CLPFGGLVRPLLSSLSYSGREK
	Ť		1			VQLPLINLPP*SGIDGGFLAHVG
						\NHGSFPWKTFPFEGRVNSGPK
						GLLAVILRPGTRHLVCLANVLS
			1			KLGILNHVPRPCSGRHGCPCSLP
						SFFLPMQIVKCNEMYQDILEIKT
21353	51721	A	21481	1	1068	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
21354	51722	Α	21482	955	2548	MSLCGARANAKMMAAYNGGT
						SAAAAGHHHHHHHHLPHLPPP
						HLHHHHHPQHHLHPGSAAAVH
						PVQQHTSSAAAAAAAAAAAAA
						MLNPGQQQPYFPSPAPGQAPGP
						AAAAPAQVQAAAAATVKAHH
					:	HQHSHHPQQQLDIEPDRPIGYG
1						AFGVVWSVTDPRDGKRVALKK
						MPNVFQNLVSCKRVFRELKML
						CFFKHDNVLSALDIL*PPHIDYF
						EEIYVVTELMQSDLHKIIVSPQP
						LSSDHVKVFLYQILRGL\KYLHF
		ĺ				RLGILH\RDIKPGNLLVNSNCVL
						KICDFGLARVEELDESRHMTQE
						VVTQYYRAPEILMGSRHYSNAI
						DIWSVGCIFAELLGRRILFQAQS
						PIQQLDLITDLLGTPSLEAMRTA
						CEGAKAHILRGPHKQPSLPVLY
						TLSSQAT\H*TGHLLFAECGFDP
		1		:		SKRISAKDALSHPYLDEGRLRY
						HTCMCKCCLSPS\TGRVNT\SDL
						EPVT\NPKFDDTFEKNLSPVRQV
						KEIIHQFILEQ\QKGNRV\PLWIN
İ		1				PQSAVFKSFISSTVAQPSEMPP\S
						PLVWE
21355	51723	Α	21483	119	692	SYKSRLARKS\TGGKAP\RKQLA
						YKKPASQECGPLTGGVKKPHR
						YR\PGTVAAP*NLDGYQKSTE\L
						MIRK\LRFQR\LVREIAQ\DF\KT
						DLRFQSAVIRWLCRMAS\EAYL
						VWPFLKDTNLVCLSMPKRVTIY
						AKKTFQLSTPASRGERALRIHS
						WMGKHFILKKKKKKFLFFLLL
						VVLNVRYFFFHGVKRYLSI
21356	51724	Α	21484	2	363	RFTKVEMKEK\MLSAAREKGR
						VTHKGKPISLKADLSAETLQAR
1						REWGPIFNILKEKNFQPRISYPA
						KLSFISEQEIKYFTDKQMLRDFV
						ITRPALKELLKEALNMERNNW
						YQPLQKHAKL

SEQ ID NO:	SEQ ID NO: of peptide sequence	ı	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21357	51725	A	21485		1468	MRNLRNIQKAKATINLGISFDE GLKGFMWSRIEAAPLPNTRWC LPHGYSGPANSGGCYYITTGMS CWLSDLYPRHDGVHVGLQGK DLEHDIPVDLGSRPTSVDPVSRS APGPQGPKAPVDSGSRTTPANP GAGPSPVDPSSRPAPVASGFRLI MIDQNTKLIPVPGWPLRLRLKI YLSSRSASLDSGSRFAPVDTNFI PVPTDRGHSHAPEDPVDRLTSV DPTKEKNWMENEFDKLTEVGF RRSVITNFSELKEHVLTHRKEA KNLEKRLDEWLSRITCAEKSLN DLMELKTTVRELREAYTSFMS QFDQAEERISVIEDQINEIKLKSL QAQEDARHRNTDQRSSENRRS EPWSLEERKREQWNSLKQNAD QQDTEAMSDYK/R/YSSET*RRR *LFCLLRKVHSKEGP*KKDQRE TRNQKGLGFLRDRPQWVSGKM KQECLQLGTPWP*IIIGQWQYFP RDGMGVRTDFCRIFQRLARMR LCEGCLMNS

2732

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
		l		sequence		
21358	51726	Ā	21486	1	2589	MSPVTLWVRTSSRGEDRCGMC
						TPGRDTGFAPISGMCSKYRSCTI
						NEDTGLGLAFTMPMSLDTKHIP
						NFGKVHGYIMRHHKNEDTIPW
						RRNTKKEEARGTQPLEADDQN
				-		PGWEDGKPGSLENEGRRSHRG
						NPAPDMPQPPKEDLFILPDEYK
						SCLRHKRSLLRSHRNEELNVET
				}		LVVVDKKMMQNHGHENITTY
						VLTILNMERILKKTSFTIIHPSA
		ŀ		]		MERCPVLNLQIRDSLWNHCFSL
						ATTVHRFHPVGRHALKKMYKL
						PSAYINVSITKYTQRFALTYISC
				†		VFHRNAAFVAVPGEGRCCAGV
					•	DVPAPMSLRSPGPHWPQEHGA
						LWISCTVRGMASCLGQLCRVS
						MRVRANIGKVQKPRLRISNGK
}						GLHTQSELSGSAYLEKGWRGA
						APGLDSRNEKGTEGARDSGTPG
						RQSQTLQRGPPGPVPGSSKIAPQ
						TLTRQGSMTASRKLVLLLPEFC
						ERTGEQSGQLYPPSPLLVEWV
						MPAGTSSTVDPLLPELCRRWER
						QALSLCRLYRQRTSVSIKALCD
						HTETRSVALSTCQGLECNSSPA
						RKQNWMENEFEELTEVGFRKS
				ł		VITKFSKLKEHILTHRKEAKNLE
						KRLDEWLTRITSVEKSLNDLME
		İ		1		LKTTVRK\LHEAYTSFNSRFDQ
				}		VEERILVIEHQINEIKQEDKVRE
		ł				KRVKRNEQSLQEIWDCVKRPN
						LCLIGVPESDGENGTKLENT\LQ
21359	51727	Α	21487	260	1292	RLCAFNKRMTFQFNFTIEDHLE
İ						NELTPIRDGALTLDSSKELSVSE
						SQKGEERDRKCSAEQFDLPQD
						HLWEHKSMENAAPSQDTDSPL
						SAASSSRNLEPHGKQPSLRAAK
						EHAMPKDLKKMLENKVIETLP
						GFQHVKLSVVKTILLKENFPGE
		1				NIVSKSFSSHSDLITGVYEGGLK
						IWECTFDLLAYFTKAKVKFAG
						KKVLDLGCGSGLLGITAFKGGS
						KEIHFQDYNSMVIDEVT\LPNV
						VANSTLEDEENDVNEPDVKRC
						RKPKVTQLYKCRFFSGEWSEFC
						KLVLSSEKLFVKYDLI\LTSETIY
						NPDYYSNLHQTFLRLLSKNGRV
					<u>.</u>	LLASKAHVQRNSLSRL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ł	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21360	51728	I <sub>A</sub>	21488	263	1466	RLCAFNKRMTFQFNFTIEDHL\E
21300	31720	(``	21400	203	1100	NELTPIRDG\ALTL\DSSKEL\SVS
						ESQKGRKE\RDRKCSAEQF\DLP
						QDHLWGNISSMGKWQAPS\QG
						HRTSSTSVAASSSRNL\EPHGKT
			ļ			ALLAGAGQRSMAMPKRF*RRC
-						LEN*SPLET\LPGFQHVKLSVVK
						,
1						TILLKENFPGENIVSKSFSSHSDL
			Ì			ITGVYEGGLKIWECTFDLLAYF
						TKAKVKFAGKKVLDLGCGSGL
						LGITAFKGASKEIHF\QDYNSMV
						IDEVTLPNVVA\NS\TLEDGGKM
						DVKWSPDVEKMPGNPKVTQ\L
						YKC\RFFS\GEWS*VL*ALYLSSE
						K\LFVKYDPHSSPSET\IYNP\DY
						YSNL\HQTFLRLLSKNG\RVLLA
						SKAHYFGVGGGVHLFQKFVEE
						RDVFKTRILKIIDEGLKRFIIEITF
21361	51729	Α	21489	2	367	
21362	51730	Α	21490	2	376	QTYSLRRATPRHIIVGFTKVEM
1			<u> </u>			KEKVLRAA/NKPIRLTVDLSAET
			<u> </u>			LQARKEGGPIFNILKEKNFQPRI
						SYPAKLSFISEGEIKSFTDKQML
						KDFVTTRPALQELLKEALNME
		1				RNNQYQPLQKHAKW
21363	51731	A	21491	1	690	
21364 21365	51732	A	21492	1	714	
21365	51733	A	21493 21494	1	1536 765	
21367	51734	A	21494	1	972	
21368	51735 51736	A	21493	1	2670	
21369	51730	A	21496	2	1178	
21309	51738	A	21497	2	827	
21370	51739	A	21499	1	777	
21371	51740	A	21500	168	1057	IRLKNDHDGELRNKTICINRNL
21372	31740	^	21300	100	1057	AEIKKFSENYENKETMYLNLW
					}	DVAKAVLRGQFIALNARIKKLE
		1				KSQVYNLISQLKELENQEQTNP
						KASRRQEITKIRAELKEVETWK
l						TLKKINESGRGVDCSS/DASNGT
1						KPDGE*L**VDKSR/PSEARQAN
						IHIQEIHGTPQR*SLRRATPRH/II
1						VRFTKVEMKEKMLRAARKKG
		1			1	WVTHKGKPIRLTADLLAETLQ
						ARREWGPIFNILKEKNFQLRISY
						PAKLSFIREGEIKAFTEKQILRDF
						VTTRPALQELLKEALNMERND
		<u> </u>				RYQPLQKHAKL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21373	51741		21501	2	1262	COTIOTY CVEVAPARETY AUTOC
21373	51741	А	21501	3	1263	GRTIQTKGKEV\ENFEKNLEECI
	ŀ	1				TRITNTEKCLKELMELKTKARE
	,					VREECRSLRSQCDQLEERVSAM
						EDEMNEMKQEGKFREKRIKRN
						EQSLQEIWDYVKRPNLRLIGVP
ŀ	:	l		:		ESDVENGTKLENTLQDIIQENFP
						NLARQANVQIQERQRTPQRYSS
ŀ						RRATPRHIIVRFTKVEMKEKMG
İ						LLVPNWTNHSPLFRAILFDYKG
						FCRFGTTHQTGFSPAGANQRGP
		ŀ				LAATLSGPGGEGQSAVARLTGL
		ł				IGVPESDVEKETKLETTLQDIIQ
						ENFPILARQANVQIQEIQRTPQR
1	ŀ	ł				YSSRRATPRHIIVRFTKVEMKE
					:	KMLRAAREKGRVTLKGKPIRL
	ŀ					TADLSAETLQARREWGPIFNIL
		ľ	ĺ			KEKNFQPRISYPAKLSFISEGEIK
						SFTDKQMLRDFVTTRPALKELL
						KEALNMERNNRYQPLQNHAK
21374	51742	В	21502	1	2356	
21375	51743	A	21503	2	484	
21376	51744	A	21504	2	902	ASFFSKKTSGFGGWGAGGPGD
21370	] 1 / 4 4	^	21304	2	702	PGMGNHGGFRGGFSRGIRGWG
	•	1		!		HSRGQGPGQGRGAHRGKTKDK
						EWTSIPKLGHLVKDMKIKFLEEI
	·					YLFFLAIKEFEIIDFFLGASLKDE
		İ				VLKIMPVQKQTHASQRTRLKG
	ĺ				•	FIVIGDYNG/HVGLGVKCSKEV
						A/TAIRGAIILG*LSIVPMCRGY
	l					WGSKIGKPHTVPCKVTGHCGS
						VLVHFIPAPRGTGIVSVPMP/KK
						LLLMAGIYDCY/ARSCTATLGN
				·		FTKATFDAISKTYSYPTPNLWK
						ETVFTKSP/YEFRNSGQEFTDYL
						VKT/HIRVSVQRTQAPAVATT
21377	51745	A	21505	2	407	STISCGGNFPIPRGGSGVASLER
						AESSSTEPAKAIKPIDRKSVHQI
		]				CSGPVVPSLSTAVKELVENSLD
		1				AGATNIDLKLKDYGVDLIEVSG
		1				NGCGVEEENFEGLTLKHHTSKI
						QEFADL\LSDVTISTCHVSAKVG
21378	51746	Α	21506	362	490	
21379	51747	Α	21507	3009	3889	
21380	51748	Α	21508	1	504	LDDLHKWFLSSFLLTAISIGIKF
						LLKIHISRGSGVLLQLLPGNPSP
						AAHGTEELEPKFLEQGYLRAVL
						PGPRVHATA*LEPTVTDQGYAR
	]					AGLPIQEVEDVLFGVLVVLLHV
						IGQVAEDRNQVAAQGHHQES/C
						PAQCWGDCTHLEIAGWRGIRRS
				,		1
	L	<u> </u>				DANPRSQRKGFG

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21381	51749	Α	21509	2	1101	WTERRLGSVRGDRLRERSRPA VNRTPRMSRVPSGAPRGTPSRR WWMMKCWDHQPLG*IPAPHV GPLAG/IRKKEWSDESEEEPEKE LAPEPEETWVVEMLCGLKMKL KQQRVSPILPEHHKDFNSQLAP GVDPSPPHRSFCWKRKREWWD ESEESLEEEPRKVLAPEPEEIWV AEMLCGLKMKLKRRVSLVLP EHHEAFNRLLEDPVIKRFLAWD KDLRVSDKYLLAMVIAYFSRA GLPSWQYQRIHFFLALYLAND MEEDDEDPKQNIFYFLYGKTRS RIPLVRNRRFQLCRCLNPRARK NRSQIALFQKLRFQFFCSMSGR AWVSREELEENTGPRGDVDFQ QELYSNANGRQQERGEEPFVQI
21382	51750	Α	21510	3	918	
21383	51751	Α	21511	1	792	
21384	51752	Α	21512	3007	3885	
21385	51753	A	21513	615	845	GHTASPPPRSPQAQGRAPSVAP PPTTPQICPTTPSSFSSKRTYAG GWDLPQGQADC*SPCGAQGG WVTLAVASASS
21386	51754	A	21514	357	562	GSSTCGRETERVVLKKRKRSPT HFQYRNSPTAEARPQLE\PTQW GRRSHSPRPSAPLPCSCPQKSRK SV
21387	51755	Α	21515	1	1335	
21388	51756	A	21517	537	1164	SSTEPAKAIKPIDRKSVHQICSG PVVLSLSTAVKELVENSLDAGA TNIDLKLKDYGVDLIEVSDNGC GVEEENFEGLISF\TLKHHTCKI QEFADLTEVETFGFQGEALSSL CALSDVTISTCHASVKVGTRLV FDHDGKIIQETPYP\HPRGTTVS VKQLFSTLPVRHKEFQRNIKK\R ACFPFAFCRDCQFPEASP\AMLP VQPAELMAV
21389	51757	A	21517	224	903	GFTWYIGMAQPKQERVARARH QRSETARHQRSETAKTPTLGNR QTPTLGNRQTPRLGIHARPRRR ATTSLLTLLRAFGKKTQS/STEP AKAIKPIDRKSVHQICSGPVVLS LSTAVKKIVGNSLDAGATNIDL KLKDYGMDLIEVSGNGCGVEE ENFEGLSWDSTGVFDHDGKIIQ KTPYPHPRGTTVSVKQLFSTLP VRHKEFQRNIKKPVGEKYMRS VDTQACSCI

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21390	51758	A	21518	2	1321	VKSRAALLGPGA/LSPLEVSRRR PWTERRLGSVRGDRLRERSRPA VNRTPRMSRVPSGAPRGTPSRR WWMMKCWDHQPLG*IPAPHV GPLAG/IRKREWSDESEEEPEKE LAPEPEETWVVEMLCGLKMKL KQQRVSSILPEHHKDFNSQLAP GVNPSPPHRSFCWKRKMEWW DQSEESLEEEPRKVLAPEPEEIW VAEMLCGLKMKLKRRRVSLVL PEHHEAFNRLLEDPVIKRFLAW DKDLRVSDKYLLAMVIAYFSR AGFPSWQYQRIHFFLALYLAND MEEDDEDSKQNIFHFLYRKNRS RIPLLRKRWFQLGHSMNPRARK NRSRIPLLRKRRFQLYRSTNPRA RKNRSRIPLLCKRRFQLYRSTNPRA RKNRSRIPLLCKRRFQLYRSMN SRARKNRSQIVLFQKRRFHFFCS MSCRAWVSPEELEENTGPRGD VDFQOELYSNANGRHQAGGEE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21391	51759	Α	21519	1	2654	RSACGSPAWNFPSPRGGSGVAS
						ME\RAESSSTEPAKAIKPIDQKS
	1					VHQICSGQVVLSLSTAVKELVE
	Ţ	1				NSLDAGATNIDLKLKDYGVDLI
		1				EVSDNGCGVEEENFEGLTLKH
						HTSKIHEFADLTQVETFGFRGE
						ALSSLCALSDVTIFTCHASAKV
						GTRLMFDHNGKIIQKTPYPRPR
						GTTVSVQQLFSTLPVRHKEFQR
						NIKKEYAKMVQVLHAYC\SFQ
						QASV*VAPISLDK/RKRQPVVCT
						GGSPSIKENIGSVFGQKQLQSLI
	1					PFVQLPPSDSVCEEYGLSCSDA
						LHNLFYISGFISQCTHGVGRSST
						DRQFFFINRRPCDPAKVCRLVN
			ŀ			EVYHMYNRHQYPFVVLNISVD
						SECVDINVTPDKRQILLQEEKLL
						LAVLKTSLIGMFDSDVNKLNVS
j						QQPLLDVEGNLIKMHAADLEK
1			j			PMVEKQDQSPSLRTGEEKKDV
						SISRLREAFSLRHTTENKPHSPK
		1				TPEPRRSPLGQKRGMLSSSTSG
						AISDKGVLRPQKEAVSSSHGPS
						DPTDRAEVEKDSGHGSTSVDSE
						GFSIPDTGSHCSSEYAASSPGDR
	1					GSQEHVDSQEKAPETDDSFSDV
						DCHSNQEDTGCKFRVLPQPTNL
						ATPNTKRFKKEEILSSSDICQKL
						VNTQDMSASQVDVAVKINKKV
	İ					VPLDFSMSSLAKRIKQLHHEAQ
						QSEGEQNYRKFRAKICPGENQA
				İ		AEDELRKEISKTMFAEMEIIGOF
21392	51760	A	21520	238	413	
21393	51761	Α	21521	288	467	
21394	51762	Α	21522	1	3038	MDVGWDLGEDCWQEHLHTAF
		1				SFDCWPCSQNGDRVPSVNPVG
						GGAPRSYLQVASAECWAAAPA
<u> </u>		1				VHVGEPVHAGGLHTERGADPV
		1		}		IGLYLVHRGGACQTPTVGNRQ
		1				TPTLGIHARPRRRATTSLLTLLL
		1				AFGKNAVRTEPAKAIKPIDRKS
		1				VHQICSGPVVPSLSTAVKELVE
		1				NSLDAGATNIDLKLKDKESGL\I
						KVQANEWGVKEETSKGLLLKH
						HTSKIQEFADLPQVETFGFRGE
		1				ALSSLCALSDVTISTCHV
21395	51763	A	21523	34	256	ETGPPGSAPGRLEFTNIKRVVVP
[						QERVELETLLVSYFSPRTPRRTP
						PPFPHPCS*EDNVITVFSSIKNGP
						GSSR
	. L				<u> </u>	

LLVWYIGMAQPKQERVARARH QRSETARHQRSETA/KTPTLGN RQTPTLGNRQTPRLGIHARPRR RATTSLITLLRAFGKKTQS/STE PAKAIKPIDRKSVHQICSGQVVU SLSTAVKKIVENSLDAGATNID LKLKDYGMDLIEVSGNGCGVE EENFKGLTLKHHTSKIQEFADL TRVETFGFRGEALSSLCALSDV TISTCHVSAKVGTRLVFDHDGK IIQKTPYPHPRGTTVSVKQLFST LPVRHKEFQRNIKKAQR/PQAR/ CYRCS*PNVSPGPGMAASTLS A*PRSQSSYFISCKA/EDNAICPI AEQKAEGKFSSRYRNHDSAEN RRDGSMTRPRTDWTLKRACTE KRRPQHASSFEWRGHSLCWDR EQRDAETLKMLLDSGLSVQKK TKDRTETRFGEMGQILGKIMMS HQPQPQEEQSPQRSTSGYPLQE VVDDEVSGPSAPGVDPSPPRRS LGWKRKRECLDESDDEPEKEL APEPEETWVAETLCGLKMKAK RRRVSLVLPEYYEAFNRLLEDP VIKRLLAWDKDLRVSDKIPSEP TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM	SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
21396 51764 A 21524 727 2953 HVSGYDDVPLGTLSLDIWPPAL FSVVSFAAFFTPPIEAVLKSFSC LLVWYIGMAQPKQERVARARH QRSETARHQRSETAKTPTLGM RQTPTLGNRQTPRLGIHARPRR RATTSLLTLLRAFGKKTQS:STE PAKAIKPIDRKSVHQICSGQVVL SLSTAVKKIVENSLDAGATNID LKLKDYGMDLIEVSGNGCGVE EENFKGLTLKHHTSKIQEFADL TRVETFGFRGEALSSLCALSDV TISTCHVSAKVGTRLVFDHDGK IIQKTPYPHPGTTVSVKQLFST LPVRHKEFQRNIKKAQR/PQAR/ CYRCS*PNVSPGPGMAAASTLS A*PRSQSSYFISCKA/EDNAICPI AEQKAEGKFSSRYRNHDSAEN RRDGSMTRPRTDWTLKRACTE KRRPQHASSFEWRGHSLCWDR EQRDAETLKMLLDSGLSVQKK TKDRTETTFFEGMGQILGKIMMS HQPQPQEGSPORSTSGYPLQE VVDDEVSGPSAPGVDPSPPRRS LGWKRKRCLDESDDEPEKEL APEPETWVAETLCGLKMKAK RRRVSLVLPEYYEAFNRLLEDP VIKRLLAWDKDLRVSDKIPSEP TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHI PPLRELWFQLCRYMNPRARKN	NO:	1	hod		•		
21396 51764 A 21524 727 2953 HVSGYDDVPLGTLSLDIWPPAL FSVVSFAAFFFTPPIEAVLKSFSC LLVWYIGMAQPKQERVARARH QRSETARHQRSETA/KTPTLGN RQTPTLGNRQTPRLGIHARPRR RATTSLLTLLRAFGKKTQS/STE PAKAIKPIDRKSVHQICSGQVVL SLSTAVKKIVENSLDAGATNID LKLKDYGMDLIEVSGNGCGVE EENFKGLTLKHHTSKIQEFADL TRVETFGFRGEALSSLCALSDV TISTCHVSAKVGTRLVFDHDGK IIQKTPYPHPRGTTVSVKQLFST LPVRHKEFQRNIKKAQR/PQAR/ CYRCS*PNVSPGFMAAASTLS A*PRSQSSYFISCKA/EDNAICH APKAGSKTRPTDWTLKRACTE KRRPQHASSFEWRGHSLCWDR EQRDAETLKMLLDSGLSVQKK TKDRTETRFGEMGQILGKIMMS HQPQPQEEQSPQRSTSGYPLQE VVDDEVSGPSAPGVDPSPPRRS LGWKRKRECLDESDDEPEKEL APEPETWVAETLCGLKMKAK RRRVSLVLPEYYEAFNRLLEDP VIKRLLAWDKDLRVSDKIPSEP TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHI PLRELWFQLCRYMNPRARKN		sequence		09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
FSVVSFAAFFTPPIEAVLKSFSC LLVWYIGMAQPKQERVARARH QRSETARHQRSETAKTPITLGN RQTPTLGNRQTPRLGIHARPRR RATTSLTLLLAFGKKTQS/STE PAKAIKPIDRKSVHQICSGQVVL SLSTAVKKIVENSLDAGATNID LKLKDYGMDLIEVSGNGCGVE EENFKGLTLKHHTSKIQEFADL TTVETTFGFRGEALSSLCALSDV TISTCHVSAKVGTRLVFDHDGK IIQKTPYPHPRGTTVSVKQLFST LPVRHKEFQRNIKKAQR/PQAR/ CYRCS*PNVSPGPGMAAASTLS A*PRSQSYFISCKA/EDNAICPI AEQKAEGKFSSRYRNHDSAEN RRDGSMTRPRTDWTLKRACTE KRRPQHASSFEWRGHSLCWDR EQRDAETLKMLLDSGLSVQKK TKDRTETRFGEMGQILGKIMMS HQPQPQEEQSPQRSTSGYPLQE VVDDEVSGPSAPGVDPSPPRS LGWKRKRECLDESDDEPEKEL APEPETWVAETLCGLKMKAK RRRVSLVLPEYYEAFNRLLEDP VIKRLLAWDKDLRVSDKIPSEP TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHIP PLRELWFQLCRYMNPRARKN					sequence		
FSVVSFAAFFTPPIEAVLKSFSC LLVWYIGMAQPKQERVARARH QRSETARHQRSETAKTPITLGN RQTPTLGNRQTPRLGIHARPRR RATTSLTLLLAFGKKTQS/STE PAKAIKPIDRKSVHQICSGQVVL SLSTAVKKIVENSLDAGATNID LKLKDYGMDLIEVSGNGCGVE EENFKGLTLKHHTSKIQEFADL TTVETTFGFRGEALSSLCALSDV TISTCHVSAKVGTRLVFDHDGK IIQKTPYPHPRGTTVSVKQLFST LPVRHKEFQRNIKKAQR/PQAR/ CYRCS*PNVSPGPGMAAASTLS A*PRSQSYFISCKA/EDNAICPI AEQKAEGKFSSRYRNHDSAEN RRDGSMTRPRTDWTLKRACTE KRRPQHASSFEWRGHSLCWDR EQRDAETLKMLLDSGLSVQKK TKDRTETRFGEMGQILGKIMMS HQPQPQEEQSPQRSTSGYPLQE VVDDEVSGPSAPGVDPSPPRS LGWKRKRECLDESDDEPEKEL APEPETWVAETLCGLKMKAK RRRVSLVLPEYYEAFNRLLEDP VIKRLLAWDKDLRVSDKIPSEP TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHIP PLRELWFQLCRYMNPRARKN	21396	51764	A	21524	727	2953	HVSGYDDVPLGTLSLDIWPPAL
QRSETARHQRSETA/KTPTLGN RQTPTLGNRQTPRLGIHARPRR RATTSLLTLLRAFGKKTQS/STE PAKAIKPIDRKSVHQICSQQVVL SLSTAVKKIVENSLDAGATNID LKLKDYGMDLIEVSGNGCGVE EENFKGLTLKHHTSKIQEFADL TRVETFGFRGEALSSLCALSDV TISTCHVSAKVGTRLVFDHDGK IIQKTPYPHPRGTTVSVKQLFST LPVRHKEFQRNIKKAQR/PQAR/ CYYRCS*PNVSPGFGMAAASTLS A*PRSQSSYFISCKA/EDNAICPI AEQKAEGKFSSRYRNHDSAEN RRDGSMTRPRTDWTLKRACTE KRRPQHASSFEWRGHSLCWDR EQRDAETLKMLLDSGLSVQKK TKDRTETRFGEMGQILGKIMMS HQPPQEEQSPQRSTSGYPLQE VVDDEVSGPSAPGVDPSPPRRS LGWKRKRECLDESDDEPEKEL APEPEETWVAETLCGLKMKAK RRRVSLVLPEYYEAFNRLLEDP VIKRLLAWDKDLRVSGLISCH			1		/		FSVVSFAAFFFTPPIEAVLKSFSC
QRSETARHQRSETA/KTPTLGN RQTPTLGNRQTPRLGIHARPRR RATTSLLTLLRAFGKKTQS/STE PAKAIKPIDRKSVHQICSQQVVL SLSTAVKKIVENSLDAGATNID LKLKDYGMDLIEVSGNGCGVE EENFKGLTLKHHTSKIQEFADL TRVETFGFRGEALSSLCALSDV TISTCHVSAKVGTRLVFDHDGK IIQKTPYPHPRGTTVSVKQLFST LPVRHKEFQRNIKKAQR/PQAR/ CYYRCS*PNVSPGFGMAAASTLS A*PRSQSSYFISCKA/EDNAICPI AEQKAEGKFSSRYRNHDSAEN RRDGSMTRPRTDWTLKRACTE KRRPQHASSFEWRGHSLCWDR EQRDAETLKMLLDSGLSVQKK TKDRTETRFGEMGQILGKIMMS HQPPQEEQSPQRSTSGYPLQE VVDDEVSGPSAPGVDPSPPRRS LGWKRKRECLDESDDEPEKEL APEPEETWVAETLCGLKMKAK RRRVSLVLPEYYEAFNRLLEDP VIKRLLAWDKDLRVSGLISCH							LLVWYIGMAQPKQERVARARH
RQTPTLGNRQTPRLGIHARPRR RATTSLLTLLRAFGKKTQS/STE PAKAIKPIDRKSVHQICSGQVVI SLSTAVKKIVENSLDAGATNID LKLKDYGMDLIEVSGNGCGVE EENFKGLTLKHHTSKIQEFADL TRVETFGFRGEALSSLCALSDV TISTCHVSAKVGTRLVFDHDGK IIQKTPYPHPRGTTVSVKQLFST LPVRHKEFQRNIKKAQR/PQAR/ CYRCS*PNVSPGPGMAAASTLS A*PRSQSSYFISCKA/EDNAICPI AEQKAEGKFSSRYRNHDSAEN RRDGSMTRPRTDWTLKRACTE KRRPQHASSFEWRGHSLCWDR EQRDAETLKMLDSGLSVQKK TKDRTETFFGEMGQILGKIMMS HQPQPQEEQSPQRSTSGYPLQE VVDDEVSGPSAPGVDPSPPRRS LGWKRKRECLDESDDEPEKEL APEPEETWVAETLCGLKMKAK RRRVSLVLPEYYEAFNRLLEDP VIKRLLAWDKDLRVSDKIPSEP TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHI PLLRELWFQLCRYMNPRARKN							
RATTSLLTLLRAFGKKTQS/STE PAKAIKPIDRKSVHQICSGQVVL SLSTAVKKIVENSLDAGATNID LKLKDYGMDLIEVSGNGCGVE EENFKGLTLKHHTSKIQEFADL TRVETFGFRGEALSSLCALSDV TISTCHVSAKVGTRLVFDHDGK IIOKTPYPHPRGTTVSVKQLFST LPVRHKEFQRNIKKAQR/PQAR/ CYRCS*PNVSPGPGMAAASTLS A*PRSQSSYFISCKA/EDNAICPI AEQKAEGKFSSRYRNHDSAEN RRDGSMTRPRTDWTLKRACTE KRRPQHASSFEWRGHSLCWDR EQRDAETLKMLLDSGLSVQKK TKDRTETRFGEMGQILGKIMMS HQPQPQEEQSPQRSTSGYPLQE VVDDEVSGPSAPGVDPSPPRRS LGWKRKRECLDESDDEPEKEL APEPEETWVAETLCGLKMKAK RRRVSLVLPEYYEAFNRLLEDP VIKRLAWDKDLRVSDKIPSEP TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHI PLLRELWFQLCRYMNPRARKN					5		1 -
SLSTAVKKIVENSLDAGATNID LKLKDYGMDLIEVSGNGCGVE EENFKGLTLKHHTSKIQEFADL TRVETFGFRGEALSSLCALSDV TISTCHVSAKVGTRLVFDHDGK IIQKTPYPHPRGTTVSVKQLFST LPVRHKEFQRNIKKAQR/PQAR/ CYRCS*PNVSPGPGMAAASTLS A*PRSQSSYFISCKA/EDNAICPI AEQKAEGKFSSRYRNHDSAEN RRDGSMTRPRTDWTLKRACTE KRRPQHASSFEWRGHSLCWDR EQRDAETLKMLLDSGLSVQKK TKDRTETRFGEMGQILGKIMMS HQPQPQEEQSPQRSTSGYPLQE VVDDEVSGPSAPGVDPSPPRRS LGWKRKRECLDESDDEPEKEL APEPEETWVAETLCGLKMKAK RRRVSLVLPEYYEAFNRLLEDP VIKRLLAWDKDLRVSDKIPSEP TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHI PLLRELWFQLCRYMNPRARKN	•						RATTSLLTLLRAFGKKTQS/STE
LKLKDYGMDLIEVSGNGCGVE EENFKGLTLKHHTSKIQEFADL TRVETFGFRGEALSSLCALSDV TISTCHVSAKVGTRLVFDHDGK IIQKTPYPHPRGTTVSVKQLFST LPVRHKEFQRNIKKAQR/PQAR/ CYRCS*PNVSPGPGMAAASTLS A*PRSQSSYFISCKA/EDNAICPI AEQKAEGKFSSRYRNHDSAEN RRDGSMTRPRTDWTLKRACTE KRRPQHASSFEWRGHSLCWDR EQRDAETLKMLLDSGLSVQKK TKDRTETRFGEMGQILGKIMMS HQPQPQEEQSPQRSTSGYPLQE VVDDEVSGPSAPGVDPSPPRRS LGWKRKECLDESDDEPEKEL APEPETWVAETLCGLKMKAK RRRVSLVLPEYYEAFNRLLEDP VIKRLLAWDKDLRVSDKIPSEP TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHI PLLRELWFQLCRYMNPRARKN							PAKAIKPIDRKSVHQICSGQVVL
EENFKGLTLKHHTSKIQEFADL TRVETFGFRGEALSSLCALSDV TISTCHVSAKVGTRLVFDHDGK IIQKTPYPHPRGTTVSVKQLFST LPVRHKEFQRNIKKAQR/PQAR/ CYRCS*PNVSPGPGMAAASTLS A*PRSQSSYFISCKA/EDNAICPI AEQKAEGKFSSRYRNHDSAEN RRDGSMTRPRTDWTLKRACTE KRRPQHASSFEWRGHSLCWDR EQRDAETLKMLLDSGLSVQKK TKDRTETRFGEMGQILGKIMMS HQPQPQEEQSPQRSTSGYPLQE VVDDEVSGPSAPGVDPSPPRRS LGWKRKRCLDESDDEPEKEL APEPEETWAETLCGLKMKAK RRRVSLVLPEYYEAFNRLLEDP VIKRLLAWDKDLRVSDKIPSEP TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHI PLLRELWFQLCRYMNPRARKN			ŀ				SLSTAVKKIVENSLDAGATNID
TRVETFGFRGEALSSLCALSDV TISTCHVSAKVGTRLVFDHDGK IIQKTPYPHPRGTTVSVKQLFST LPVRHKEFQRNIKKAQR/PQAR/ CYRCS*PNVSPGPGMAAASTLS A*PRSQSSYFISCKA/EDNAICPI AEQKAEGKFSSRYRNHDSAEN RRDGSMTRPRTDWTLKRACTE KRRPQHASSFEWRGHSLCWDR EQRDAETLKMLLDSGLSVQKK TKDRTETRFGEMGQILGKIMMS HQPQPQEEQSPQRSTSGYPLQE VVDDEVSGPSAPGVDPSPPRRS LGWKRKRECLDESDDEPEKEL APEPEETWVAETLCGLKMKAK RRRVSLVLPEYYEAFNRLLEDP VIKRLLAWDKDLRVSDKIPSEP TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHI PLLRELWFQLCRYMNPRARKN							LKLKDYGMDLIEVSGNGCGVE
TISTCHVSAKVGTRLVFDHDGK IIQKTPYPHPRGTTVSVKQLFST LPVRHKEFQRNIKKAQR/PQAR/ CYRCS*PNVSPGPGMAAASTLS A*PRSQSSYFISCKA/EDNAICPI AEQKAEGKFSSRYRNHDSAEN RRDGSMTRPRTDWTLKRACTE KRRPQHASSFEWRGHSLCWDR EQRDAETLKMLLDSGLSVQKK TKDRTETRFGEMGQILGKIMMS HQPQPQEEQSPQRSTSGYPLQE VVDDEVSGPSAPGVDPSPPRRS LGWKRKRECLDESDDEPEKEL APEPEETWVAETLCGLKMKAK RRRVSLVLPEYYEAFNRLLEDP VIKRLLAWDKDLRVSDKIPSEP TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHI PLLRELWFQLCRYMNPRARKN							EENFKGLTLKHHTSKIQEFADL
IIQKTPYPHPRGTTVSVKQLFST LPVRHKEFQRNIKKAQR/PQAR/ CYRCS*PNVSPGPGMAAASTLS A*PRSQSSYFISCKA/EDNAICPI AEQKAEGKFSSRYRNHDSAEN RRDGSMTRPRTDWTLKRACTE KRRPQHASSFEWRGHSLCWDR EQRDAETLKMLLDSGLSVQKK TKDRTETRFGEMGQILGKIMMS HQPQPQEEQSPQRSTSGYPLQE VVDDEVSGPSAPGVDPSPPRRS LGWKRKRECLDESDDEPEKEL APEPEETWVAETLCGLKMKAK RRRVSLVLPEYYEAFNRLLEDP VIKRLLAWDKDLRVSDKIPSEP TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHI PLLRELWFQLCRYMNPRARKN							TRVETFGFRGEALSSLCALSDV
LPVRHKEFQRNIKKAQR/PQAR/ CYRCS*PNVSPGPGMAAASTLS A*PRSQSSYFISCKA/EDNAICPI AEQKAEGKFSSRYRNHDSAEN RRDGSMTRPRTDWTLKRACTE KRRPQHASSFEWRGHSLCWDR EQRDAETLKMLLDSGLSVQKK TKDRTETRFGEMGQILGKIMMS HQPQPQEEQSPQRSTSGYPLQE VVDDEVSGPSAPGVDPSPPRRS LGWKRKRECLDESDDEPEKEL APEPEETWVAETLCGLKMKAK RRRVSLVLPEYYEAFNRLLEDP VIKRLLAWDKDLRVSDKIPSEP TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHI PLLRELWFQLCRYMNPRARKN							TISTCHVSAKVGTRLVFDHDGK
CYRCS*PNVSPGPGMAAASTLS A*PRSQSSYFISCKA/EDNAICPI AEQKAEGKFSSRYRNHDSAEN RRDGSMTRPRTDWTLKRACTE KRRPQHASSFEWRGHSLCWDR EQRDAETLKMLLDSGLSVQKK TKDRTETRFGEMGQILGKIMMS HQPQPQEEQSPQRSTSGYPLQE VVDDEVSGPSAPGVDPSPPRRS LGWKRKRECLDESDDEPEKEL APEPEETWVAETLCGLKMKAK RRRVSLVLPEYYEAFNRLLEDP VIKRLLAWDKDLRVSDKIPSEP TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHI PLLRELWFQLCRYMNPRARKN			1				IIQKTPYPHPRGTTVSVKQLFST
A*PRSQSSYFISCKA/EDNAICPI AEQKAEGKFSSRYRNHDSAEN RRDGSMTRPRTDWTLKRACTE KRRPQHASSFEWRGHSLCWDR EQRDAETLKMLLDSGLSVQKK TKDRTETRFGEMGQILGKIMMS HQPQPQEEQSPQRSTSGYPLQE VVDDEVSGPSAPGVDPSPPRRS LGWKRKRECLDESDDEPEKEL APEPEETWVAETLCGLKMKAK RRRVSLVLPEYYEAFNRLLEDP VIKRLLAWDKDLRVSDKIPSEP TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHI PLLRELWFQLCRYMNPRARKN		}					LPVRHKEFQRNIKKAQR/PQAR/
AEQKAEGKFSSRYRNHDSAEN RRDGSMTRPRTDWTLKRACTE KRRPQHASSFEWRGHSLCWDR EQRDAETLKMLLDSGLSVQKK TKDRTETRFGEMGQILGKIMMS HQPQPQEEQSPQRSTSGYPLQE VVDDEVSGPSAPGVDPSPPRRS LGWKRKRECLDESDDEPEKEL APEPEETWVAETLCGLKMKAK RRRVSLVLPEYYEAFNRLLEDP VIKRLLAWDKDLRVSDKIPSEP TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHI PLLRELWFQLCRYMNPRARKN							CYRCS*PNVSPGPGMAAASTLS
RRDGSMTRPRTDWTLKRACTE KRRPQHASSFEWRGHSLCWDR EQRDAETLKMLLDSGLSVQKK TKDRTETRFGEMGQILGKIMMS HQPQPQEEQSPQRSTSGYPLQE VVDDEVSGPSAPGVDPSPPRRS LGWKRKRECLDESDDEPEKEL APEPEETWVAETLCGLKMKAK RRRVSLVLPEYYEAFNRLLEDP VIKRLLAWDKDLRVSDKIPSEP TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHI PLLRELWFQLCRYMNPRARKN							A*PRSQSSYFISCKA/EDNAICPI
KRRPQHASSFEWRGHSLCWDR EQRDAETLKMLLDSGLSVQKK TKDRTETRFGEMGQILGKIMMS HQPQPQEEQSPQRSTSGYPLQE VVDDEVSGPSAPGVDPSPPRRS LGWKRKRECLDESDDEPEKEL APEPEETWVAETLCGLKMKAK RRRVSLVLPEYYEAFNRLLEDP VIKRLLAWDKDLRVSDKIPSEP TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHI PLLRELWFQLCRYMNPRARKN		1					AEQKAEGKFSSRYRNHDSAEN
EQRDAETLKMLLDSGLSVQKK TKDRTETRFGEMGQILGKIMMS HQPQPQEEQSPQRSTSGYPLQE VVDDEVSGPSAPGVDPSPPRRS LGWKRKRECLDESDDEPEKEL APEPEETWVAETLCGLKMKAK RRRVSLVLPEYYEAFNRLLEDP VIKRLLAWDKDLRVSDKIPSEP TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHI PLLRELWFQLCRYMNPRARKN							RRDGSMTRPRTDWTLKRACTE
TKDRTETRFGEMGQILGKIMMS HQPQPQEEQSPQRSTSGYPLQE VVDDEVSGPSAPGVDPSPPRRS LGWKRKRECLDESDDEPEKEL APEPEETWVAETLCGLKMKAK RRRVSLVLPEYYEAFNRLLEDP VIKRLLAWDKDLRVSDKIPSEP TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHI PLLRELWFQLCRYMNPRARKN							KRRPQHASSFEWRGHSLCWDR
HQPQPQEEQSPQRSTSGYPLQE VVDDEVSGPSAPGVDPSPPRRS LGWKRKRECLDESDDEPEKEL APEPEETWVAETLCGLKMKAK RRRVSLVLPEYYEAFNRLLEDP VIKRLLAWDKDLRVSDKIPSEP TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHI PLLRELWFQLCRYMNPRARKN							EQRDAETLKMLLDSGLSVQKK
VVDDEVSGPSAPGVDPSPPRRS LGWKRKRECLDESDDEPEKEL APEPEETWVAETLCGLKMKAK RRRVSLVLPEYYEAFNRLLEDP VIKRLLAWDKDLRVSDKIPSEP TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHI PLLRELWFQLCRYMNPRARKN							TKDRTETRFGEMGQILGKIMMS
LGWKRKRECLDESDDEPEKEL APEPEETWVAETLCGLKMKAK RRRVSLVLPEYYEAFNRLLEDP VIKRLLAWDKDLRVSDKIPSEP TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHI PLLRELWFQLCRYMNPRARKN	:						HQPQPQEEQSPQRSTSGYPLQE
APEPEETWVAETLCGLKMKAK RRRVSLVLPEYYEAFNRLLEDP VIKRLLAWDKDLRVSDKIPSEP TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHI PLLRELWFQLCRYMNPRARKN							VVDDEVSGPSAPGVDPSPPRRS
RRRVSLVLPEYYEAFNRLLEDP VIKRLLAWDKDLRVSDKIPSEP TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHI PLLRELWFQLCRYMNPRARKN							LGWKRKRECLDESDDEPEKEL
VIKRLLAWDKDLRVSDKIPSEP TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHI PLLRELWFQLCRYMNPRARKN			İ				APEPEETWVAETLCGLKMKAK
TILGASPKTLPPASRICIRPSNTP PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHI PLLRELWFQLCRYMNPRARKN							RRRVSLVLPEYYEAFNRLLEDP
PPRNFHMSTVTPMLSYLANDM EEDDEAPKQNIFYFLYEETRSHI PLLRELWFQLCRYMNPRARKN			1				VIKRLLAWDKDLRVSDKIPSEP
EEDDEAPKQNIFYFLYEETRSHI PLLRELWFQLCRYMNPRARKN							TILGASPKTLPPASRICIRPSNTP
PLLRELWFQLCRYMNPRARKN			ŀ			1	PPRNFHMSTVTPMLSYLANDM
							EEDDEAPKQNIFYFLYEETRSHI
CSQIALFRKYRFHFFCSMRCRA							PLLRELWFQLCRYMNPRARKN
							CSQIALFRKYRFHFFCSMRCRA

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		•
21397	51765	A	21525	1054	2349	SCPQIIFLSSCLQVASAECWAAA PAVHVGEPVHAGGLHTERGAD
						PVIGLYLVHRGGACQTPT\IGSR
						OTPTLGIHARPRRRATTSLLTLL
						LAFGKNAVRCALIGPGSLTSRT
						RPLTEPLGEKERREVFFPPRPER
						VEHNVESSRWEPRRRGACGSR
						GGNFP\SPRGG\SGVASLERAES
						SSTEPAKAIKPIDRKSVHQICSG
,						PVVPSLRPNAVKELVENSLDAG
						A\TNVDLKLKDYGVDLIEVSG\
1						NGCGVEEENFEGL/TSLSALKH
						HTSKIREFADLTRVETFGFRGE/
						AL*ASLCALSDVTISTCHVSAK\
						VGTLTGVFDHDGKIIPENPPNPH
						PRGTTVSVKQLFSTLPV\RHKEF
						QRNIKKKRA\CFPFAFCRDCQFP
						EASPAMLPVQPAELTPR\STP\PH
						PCSLEDNVITVFSS\VKNGPGSS
21398	51766	A	21526	2	393	KMICTNGSSPSS
21399	51767	A	21527	2	165	
21400	51768	A	21528	3	196	
21401	51769	Α	21529	1	411	
21402	51770	A	21530	1	1727	GPQGVGRMAEGKAGGAAGLF
			·			AKQVQKKFSRAQEKVLQKLGK
						AVETKDERFEQSASNFYQQQA
		1				EGHKLYKDLKNFLSAVKVMHE
		1				SSKRVSETLQFIYY\SEWYGHEE
ŀ						LKAIVWNNDLLWEDYEEKL\A DQAVRTMEIYVAQFSEIKE\RIA
ŀ		1				KRGRKLVDYDSARHHLEAVQN
ŀ						AKKKDEAKTAKAEEEFNKAQT
		-				VFEDLNQELLEELPILYNSRIGC
ŀ					]	YVTIFQNISNLRDVFYREMSKL
						NHNLYEVMSKLEKQHSNKVFV
						VKGLSSSSRRSLVISPGVRTATV
						SSPLTSPTSPSTLSLKSESESVSA
			ŀ			TEDLAPDAAQGEDNSEIKELLE
			ŀ			EEEIEKEGSEAGSSEEDDPLPAC
						NGPAQAQPSPTTERAK\SQEEV
						LPSSTTPSPGGALSPSGQLSSSA
						TEVVLRTRTASEGSE\RPKKTAS
						IQRTSAPPRRPPPPRATASPRPCS
						GNIPSSPTASGGGSA\TSPRASL
						GTGTASPRTSLEI\*PNPE\PPEKP
						VRTPEAKENENIHNQNPEELCT
						SPTLMTSQVASEPGEAKKMED
						KEKDNKLISADSSEGQDQLQVS
21402	61771	<u> </u>	21521	1	1520	MVPENNNLTAPEPQEEVSTSEN
21403	51771	Α	21531	1	1530	

SEQ ID			SEQ ID NO:			Amino acid sequence ( X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	Sequence		057540,217	sequence	or populae sequence	possible nucleonal institution,
21404	51772	Α	21532	499	1587	HPRDPEAPSGPLWSASLPALYS
						QGHANEFDGSNSTSHAVLVLV
						QKHIKAVLSNKLCLSTSSLVQG
İ						VNAHLGTLIWPQPRGYLSPRSA
						IPWFSVPTVTSGLHFPWEVQ*LF
İ						LLSCWGKAHQSLPQRMPPLLC
						CQGMWVPEGSMATVGLSQQLF
						DSALLLLQKAGALNLDITGQLR
	! :					SDDNLLNTSALGRLIPEVARQF
1						PEPMPVVLKVRLGATPVAHGS
						TQTTATLRPASPFVGGSWATGP
						PNSAFPSPSSP/CLVVVNLRTPAS
						L/SSKGEALREHVCAWWMSQL
						TVASSNVGFIDTDQVRTLMGTV
						FEKPLLDHLNGSLGHGELPSLV
						WVNLHYVAPEIFVYEGYVVISS
						GTLLPEPEAKTTGRP
21405	51773	Α	21533	1385	3111	IAPSEVKEHKMVLEMLNPIHYN
						ITSIVPEAMPAATMPVLLLTGLF
						LLVWNYEGTSSIPGPGYCMGIG
						PLISHGRFLW\MGIG\SACNYYN
						RVYGEFMRVWISGEETLIISKSS
						SMFHIMKHNHYSSRFGSKLGLQ
						CIGMHEKGIIF\NNNPELWKTTR
						PFLYERLCHAPASVRMVTVCA
		1				ESLKTHLDRL\EEVTT\ESGLCG
			:		ļ	TCLTLLRRVHGWDTSNTVFL\G
						IPL\EESAIVVKIQG\YFDAWQAL LIKPDIFFKISWLYKKYE\KSVK
						DL\KDA\IEVLIAEKRRRISTEEK
					}	LEECMDFATELILAEKRGDLTR
						ENVNQCILEMLIAAPDTMSVSL
						FFMLFLIAKHP\NVEEAIIKEI\QT
						VIGERDIKIDDIQKLKVMENFIY
						ESMRYQPVVDLVMAQSL*EG*
						C*SMGYPV\KKGTNIYSWNIG\R
						VHRL*FFPKPNEFTLENFAKNV
						PYRYFQPFGFGPRGCAGKY/LS
						AMVMMKAIL\VTLLRTDST*RT
						LPGTSVFESIQEGYTTCPWHPD\
	]					ETKNML\EMDLYPKETSDRCLG
						TLRRKAGQYLTLGAFSPSVCSP
						TNHPILCPLVFILHSGTLSGPMA
21406	51774	A	21534	129	378	IVCKFSVSTHKNIRAKKPDERLP
121700		<b> </b>	21007	1.27	1	ECGTQIRKDTLKRVRRTVLHYL
						C\PPPSVPEIPAWR\GDCSP/VGK
						EGKCTPDLNPNIELRPV
L	J		<u> </u>		I	EGROTI DEITI MEDITA V

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence	liou.	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
21407	51775	A	21535	1	474	MQKQIISVTEVEWYPKYNSNM KEYWSTQNHQFEQLSTHKNIR
						AKKPDERLPECGTQIRKDTLKR PCPLDRPLWPHITAEQGSKPST
						WPAPCDP/IF*ANIHLPSLHTGPC
						RSKVQAHSSRPGCQASSTCVPR
						LQDLPC*PRFRLTLVDQDARPA
						APVYQGSRTYPADPDSRPAPVD
		ļ				SGCIISGQADW
21408	51776	A	21536	2	325	WAY BOX TO DE TO LIVE DA COROLLO
21409	51777	Α	21537	2	1011	WAFSITERREFQVYYNSCPSSS
						WIGDARAFSPPDPTGLYPIYSQ
		ĺ				DTQTFRLWLNYNTDFPGSPAC
						KWQAMELLSPQSPFRVDRGEH LPFLVKGARYTLVPAGQEGAL
						AAWLEALRGQLGRRGAVVSM
						MDAEGLERSSPDCAMGLSDGE
						WQLVLN\VWGKVEADIPGHG\Q
						EVLIRLFKGHPETL\EKFDK/FTR
						HLKSED\EMKASED\LKKHMCH
						TVLTP\LGVHILRE\KGHH*RQR
						FKPLGTSSHCHPSTKIPPLKVTL
						EFHLPGIAFIQVLAEQAIPGDFG\
						AECPGAQLKQRPWELF\RKNNG
						LPTYKEAGAFQGARALAGFPTP
						THLGPPGFKKRAGV
21410	51778	Α	21538	157	322	
21411	51779	В	21539	1	975	
21412	51780	Α	21540	218	388	RKLPLAGLPCINRQHS*SNVNIC
						FLTPV*KIQN*LQSEQFLPYPTIV
21412	51701	<u> </u>	21541	((2)	022	IITLCFFAV
21413	51781	Α	21541	663	933	GCAGPÜLVNQMFGKQYYLGKS HHP\SLVSLNRGAHCTVGKPQG
						PSGP*KLGYYFKVFPPCG*FWK
						YGPSWGMEERAWTGSPLGPGP
					]	PLKKV
21414	51782	A	21542	389	1033	NQGLRNLGLCRTCLVNKMFTS
						SILGKSHRHSLVSINQGNNALW
						KAAG\PLPLKAGYC\QGFSPCHS
					İ	LKYG\SWDEKDLTVPQPDTLKG
						SVLRWISKRGKPLAVEIEEGHC
						L\CLPLGTECLGIK/PPIVHLFNSE
						IGE\NRPYGGR*DMFCSNGCLG*
						FL*LPLRCLG\GEKHKSGL/HVHI
						PVIVPPLELNYDIDSFAHMFF\V
						DLS\LIITLLSCYIPFC
21415	51783	A	21543	82	163	
21416	51784	В	21544	1	211	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21417	51785	Α	21545	71	1086	NMNLSLVLAAFCLGIASAVPKF
						DQNLDTKWYQWKATHRRLYG
						ANEEGWRRAVWEKNMKMIEL
						HNG/EKYSQGKH\GFTMAMNA
						FGDMTNEEFRQMMGCFRNQKF
						RKGKVFREPLFLDLPKSVDWR
						KKGYVTPVKNQKQCGSCWAFS
						ATGALEGQMFRKTGKLVSLSE
						QNLVDCSRPQGNQGCNGGFMA
						RAFQYVKENGGLDSEESYPYV
						AVDEICKYRPENSVANDTGFTV
						VAPGKEKALMKAVATVGPISV
						AMDAGHSSFQFYKSGI\YF*PDC
						ISKNLDQGVLVVGYVFEGANS
						NNSKYWLVKNSWGP\EWGS\N
						GYVKIAKD\KNNHCGIATAASY
21418	51786	A	21546	170	412	LWLSFLLMHWH*IPLAAVSVFT
						VSN/PILDFS*IQLKTPKHFGNLG
						NTGFLGKQLRLLLRSLVSVSRL
						HCLLGYRRTQMPPL
21419	51787	Α	21547	1	753	
21420	51788	Α	21548	125	464	TAVFLTQSLFGGLFTRTRMKFG
						TMTRI/RGDLPWEINPLSSCSLL
						HEKDPPTTSGPQTNQPKKHLTN
						FKSRKVERTKGLLKTHLTKLSH
						QLKKAWTILLPLSLLRIQACPR
						NAT
21421	51789	A	21549	1	753	
21422	51790	A	21550	291	671	
21423	51791	Α	21551	10	453	TALLLTQSLFGGLFTRTRMKFG
						AMTRIG\DLPWEINPLSSCSLLC
						EKHPPTTSGPQTDQPKKHLTNF
						KSETKETCFIHEPKTLAPVTDW
						EGSLPLVFNHCRDGLSDHSATF
						QGCQTMQGCLPWSFTLSGKSR
						FPGAGASTPQPLLLHP
21424	51792	A	21552	740	1010	TAMLLKQSLFGCLFTRTRMKF
						GAVTRIG\DLP*ETNPLSSCSLLR
	1					DKDPPTTSSPQTHQPKKHLTNF
						KSGCSSPGRARSQSFLSLCSSTL
21425	51793	A	21553	134	358	TAMLLTQSLFGGLFTRTRMKFG
						AVIRIG\DLLWEINPLSSCSLLHE
	}					KDPPTTSGPQTDQPKKHLTNFK
						SSCSSPG
21426	51794	A	21554	392	645	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ŀ	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21427	51795	Α	21555	1	1900	MQNALESLSNRIEQVEERNSEL
						EDKFFELTQSNKDKEKRIRKYK
						QSLKEVWDYVKGPNLRIISVPE
						EEENSKSLENIFGEIIEENLPSLA
						RHLDNQIQEAQRTLWKLIAKRS
						LPRHIVIRLSKVKTKERILRAVR
				:	:	QKHQVTHLTCKESYKLKVKG
						WKRHFMQMDTKNEQGIKNEN
						HLIISIDVEKASDKIQQRFMIKTL
						SKIGIQGKYLNVIKAIYDKPTAN
						ITLNREK/LESIPSEKLKAFPLRT
						GTRQGCPLSPLFFNTVLEVLAR
						AIRQEKEIKGIQIGKEEVKLSLF
						ADDMIMYLENPKDSSRKLLEWI
						KESNKVSGYKTHVHKSVALLY
						TKQ*\QAENQIKNRNPFAIAAKK
						KKKKKKKKKRNSRIYLTKESK
						DLYKENYKTLLKEIIDNMNKW
						KHIPCSWMGRISIVKKL*\LQKA
						IYKLNAILIKIPPLLFTELEKTILK
						FIWKSNKS\RIAKVILSKKNKSG
						GITLPDFKLYYTIRKINSKWIKD
						LNLRPETIQILEDNIGKTLLEIGL
			:			GKDFMTKNPKANAKKKTKINR
						WDLIKLKSFCTAKGTVSRVKRT
						HRVGENLPHYTSIHPTIYTIYTS
						DKGLISRIYNEFKQIVTRTTCVF
						LHPPHPKAKTTSYGDQLLGGN
21428	51796	C	21556	10	339	
21429	51797	В	21557	97	1458	
21430	51798	Α	21558	1	387	
21431	51799	Α	21559	3	604	GSVIAYYWSEFSIPQHLVEEAE
				ļ		RVMAEERVFMLPPRARSLKSFV
						VTSVVAFRESEGQGWAWDWPP
						FHGVGLGPGPLQGTGTPRAQCP
						REKTTCWPGTSL*IIRKQE*GNG
						LYQEMLYSPSYSSATDSKTVQR
	[					TQDNSCSFGLHARGVELMRFTT
	1					PGFPDSPYPAHARCQWALRGD
						ADSVLSLTFRSFDLASCDEARS
	<u> </u>					DLV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21432	51800	I IA	21560	1000	2084	FRGTPVQRGGGWADLPGRCGE
21432	31800		21500	1000	2004	LGRRLRSEEQARRVHKAPS/DF
						GTG/LKENTGKVNGLEEGVEFL
						PVNNVKKVEKHGPGRWVVLA
						AVLIGLLLVLLGIGFLVWHLQY
						RDVRVQKVFNGYMRITNENFV
						DAYENSNSTEFVSLASKVKDAL
						LYSGVPFLGPYHKESAVTAFM
1						WVWREGSVGCTPDWLGVGSG
1		ı				YSVWGTKREMVLEVLDWRAK
						EGPRCVLLEGSVIAYYWSEFSIP
ļ						QHLVEEAERVMAEERVVMLPP
						RARSLKSFVVTSVVAFPTDSKT
						VQRTQDNSCSFGLHARGVELM
						RFTTPGFPDSPYPAHARCQWAL
						RGDADSVLSLTFRSFDLASCDE
	:	ļ				RGSDLVTVYNTLSPMEPHALV
21433	51801	Α	21561	3	2820	TGALILQKGEIRVINQTTCENLL
						PQQITPRMMCVGFLSGGVDSC
						QVPPGQEGGRQFPLKETNNRK
		ŀ				GRLPKRLRPHTWCCYSLECSSY
						SLRCTEDTWTEDQKVNGLEEG
						VEFLPVNNVKKVEKHGPGRWV
		1				VLAAVLIGLLLVLLGIGFLVWH
1		1				LQYRDVRVQKVFNGYMRITNE
		1				NFVDAYENSNSTEFVSLASKVK
	1	1				DALKLLYSGVPFLGPYHKESAV
						TAFSEGSVIAYYWSEFSIPQHLV
		1				EEAERVMAEERVVMLP
21434	51802	Α	21562	111	465	
21435	51803	Α	21563	2	197	
21436	51804	Α	21564	359	405	EMQSRED*RK*YL*RIVNCVPL
		ŀ				RERLGYVCGGLGERSKEGTLWI
L _						YQGLQKSQDTGRV
21437	51805	Α	21565	208	287	IVSRSEKGWVMYVEGWEREAR
						REHYGYIKGYKNHRIHMMQSS
						KALV*LEDLRKHI*IFRCVFSNP
1						QVILML
21438	51806	С	21566	141	305	
21439	51807	A	21567	170	973	RMAACRALKAVLVDLSGTLHI
						EDAAVPG\AQEALKRLRGASVII
						RFVTNTTKESKQDLF\KRLRKL
						G\FDISEDEIFTSL\TAARSLL*\R
						KQVTPMLLVD*SGHYLIFKGIQ
						TSDPN\AVVMGIGTRTFSLSKF
						WNQAFRLLLGWSTSDSNSQSQ
						GFQEFKMGLALGPWTIL*LL*K
						YATDTKATVVGKPEKTFFLGSI
						AGHWLLNLEEAVMIG\DDCRG\
						DVGGAQDVGMLGILVKTGKYR
						ASDEEKINPPPYLTCESFPHAVD
L		1			<u></u>	

SEQ ID NO:	of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21440	51808	Α	21568	42	211	
21441	51809	Α	21569	92	366	AGSEMRPGLTWACSLARLLLSP
						SFIKHWFLESCLLSCSRSGSGYA
						QWCPPHQSASAAGHCQTPAHA
		l				QAYLSSPPQMDHSSVRSLLEAC
1						QTY*SASAAGHCQTPAHAQAY
						LSSPPQMDHSSVRSLLEACQTY
21442	51810	A	21570	1	834	
21443	51811	Α	21571	3	835	
21444	51812	Α	21572	1	1649	MPIVLVDWSDIREQKRLMVLR
		i				ASVALHGRSVTLYEKAFPLSEQ
						CSKKAHDQFLADLASILPSNTT
		l				PLIVSDAGFKVPWYKSVEKLG
						WYWLSRVRGKVQYADLGAEN
		1				WKPISNLHDMSSSHSKTLGYKR
		1				LTKSNPISCQILLYKSRSKGRKN
		ļ				QRSTRTHCHHPSPKIYSASAKEP
						WVLATNLPVEIRTPKQLVNIYS
		ŀ				KRMQIEETFRDLKSPAYGLGLR
		1				HSRTSSSERFDIMLLIALMLQLT
						CWLAGVHAQKQGWDKHFQAN
		ŀ				TVRNRNGSLLTNWPSVPPTIKE
						EENSEEELAAATTSKEQEPIGTD
						LDAVRTPEPLEEFPKREDQEGS
						PPETSLPYKWVVEAANLLIPAV
						GSSLSEALDLIESDPDAWCDLS
						KFDLPEEPSAEDSINNSLVQLQA
						SHQQQVLPPRQPSA\LVPSVTEY
						RLDGHTISDLSRSSRG\ELIPISPS
						TEVGGSGIGTPPSVLKRQRKRR
						VALSPVTENSTSLSFLDSCNSLT
				ļ		PKSTPVKTLRFS\PSQ\FLNFWN
						KQDTLELESPSLTSTPVCSQKV
						VVTTPLHRDKTPLHQKHAA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21445	51813	Α	21573	59	2384	PRPGSRSGLCRRAGERGAVRAG
						GMSRRTRCEDLDELHYQDTDS
						DVPEQRDSKCKVKWTHEEDEQ
						LRALVRQFGQQDWKFLASHFP
						NRTDQQCQYRWLRVLNPDLVK
		1				GPWTKEEDQKVIELVKKYGTK
						QWTLIAKHLKGRLGKQCRERW
						HNHLNPEVNKSCWTEEEDRIIC
						EAHKVLGNRWAEIAKMLPGRT
	1					DNAGKNHWNFTIQK/RKGDTG
	1					GFLREYKD*KPPVYLPL\ELEDK
						DGLQSAQPTEGQRSLLTNWPSV
						PPTIKEEEN\SEEELAAATTSKE
	Ì	ŀ				QEPIGTDLDAVR\TP\EPLEEFP\K
		ŀ				REDQEGSPPETD/SLPYKWVVE
		l				AVNLLIPAVVS\SLYEALDLIGS\
				İ		DPDAWCDLSRFDLPEEPSIEDSI
						NNS\LVQLQAS\HQQQILPPRQP
	]					SA\LVPSVTEYRLDGHTISDLSR
						SSRGELIPISPSTEVGG\SGIGTPP
						SVLKRQRKRRVALSPVTENSTS
					]	LSFLDSCNSLTPKSTPVKTLPFS
		l				PSQFLNFWNK\QDTLELESPLA
		Ì				DYPPQVCSQKVVVTTPLHR\DK
						TPLHQKHA\AFVTPDQKYSMD
						NTPHT\PTPFKNALEKYGPLKPL
						PQTPHL\EEDLKEVLRSEAGIELI
						IEDDIRPEKQKRKPGLRRSPIKK
						VRKSLALDIVDEDVKLMMSTL
						PKSLSCRTTAPSN\SST\LTLSGIK
	1	ŀ				EDNSL\LNQGLLAGPSPRKAAC
						VPEGPESHFHGQLAPYVPVPGK
21446	51014	_	21574	190	889	SLPRRLESPGRLAMAARGVIAP
21446	51814	A	21574	190	009	VGES/LCRYAEVPCSPSAKRPDA
						DVDQQRLVRSLIAVGQGVAAL
						_ · · · · · · · · · · · · · · · · · · ·
						AFAG\RYAL\RIWKPLE\QVITET AKGRFQTPRFSSY**GG/FLNII*
						1
						LRREA\GLILGVSP\SAGKAKIRT
						\AHR\RVMILESPQIKGGSPYVA
						AKINEAKD\LLETTTKRLMLKD
						HTEGKKNEGTSKKKKSPAKYS
						KTWSFLIFYMGFDHSLIFRPLSC
						ITNKMLIVLLFIIF

NO: of peptide sequence   Nod   in USSN   location of first   codon for least amino acid   sequence   seque	vn,
21447   51815	
21447   51815   A   21575   2   652   EGYIC/RHFTGSSALLTRT   GVIANVA VVRINSPNSIR:   CKTLQEVTQLSQEAQRIV   KSTKPIVAAINGSCLGGG   SCQYRIATKDRKTVLGTT   GALPGAGGTQRLPKMVC   LDMMLTGRSIRADRAKK   DQLVEPLGPGLKPPEERT   EVAITFAKGLADKKISPK   LVEKLTAYAMTIPFVRQU   LVEKLTAYAM	ertion)
GVIANVAVVRINSPNSIRI CKTLQEVTQLSQEAQRIV KSTKPIVAAINGSCLGGG SCQYRIATKDRKTVLGTI GALPGAGGTQRLPKMVG LDMMLTGRSIRADRAKK DQLVEPLGPGLKPPEERT EVAITFAKGLADKKISPK LVEKLTAYAMTIPFVRQG  21448 51816 C 21576 98 316  21449 51817 A 21577 35 247  21450 51818 A 21578 I 2391 LQLKMVACRAIGILSRFS RSRGYICRNFTGSSALLT YGVKGDVAADRINSPNS LSKELHSEFSEVMNEIWA RSAVLISSKPGCFIAGVAI AACKVILQUEVTQLSQEVA EKP*KSPQSLIVGCPSNGG GCPKMGGVCLPAFGHEC SIRVADRAKKMGLVD\QI GTQGLKPPEERTIEYLEE AKGLADKKISPKRDKGVI TA\YAMTIPFVRQQVYKK VVRKQTKGLYPAPLKIIDV IEQGSDAGYIVEGSQKFGE KESKALMGLYPHGQVLCI GAPQKDVKHLAILGAGL IAQVSVDKGLKTILKDAT DRGQQQVFKGLNDKVK. TSPLERDSIFSNLTGQLDY KRPDMGELKAVFEDP*S VLKGKVGSR*FPDHWYS	
GVIANVAVVRINSPNSIRI CKTLQEVTQLSQEAQRIV KSTKPIVAAINGSCLGGG SCQYRIATKDRKTVLGTI GALPGAGGTQRLPKMVG LDMMLTGRSIRADRAKK DQLVEPLGPGLKPPEERT EVAITFAKGLADKKISPK LVEKLTAYAMTIPFVRQG  21448 51816 C 21576 98 316  21449 51817 A 21577 35 247  21450 51818 A 21578 I 2391 LQLKMVACRAIGILSRFS RSRGYICRNFTGSSALLT YGVKGDVAADRINSPNS LSKELHSEFSEVMNEIWA RSAVLISSKPGCFIAGVAI AACKVILQUEVTQLSQEVA EKP*KSPQSLIVGCPSNGG GCPKMGGVCLPAFGHEC SIRVADRAKKMGLVD\QI GTQGLKPPEERTIEYLEE AKGLADKKISPKRDKGVI TA\YAMTIPFVRQQVYKK VVRKQTKGLYPAPLKIIDV IEQGSDAGYIVEGSQKFGE KESKALMGLYPHGQVLCI GAPQKDVKHLAILGAGL IAQVSVDKGLKTILKDAT DRGQQQVFKGLNDKVK. TSPLERDSIFSNLTGQLDY KRPDMGELKAVFEDP*S VLKGKVGSR*FPDHWYS	HITY
CKTLQEVTQLSQEAQRIV KSTKPIVAAINGSCLGGG SCQYRIATKRKTVLGTI GALPGAGTQRLPKMVG LDMMLTGRSIRADRAKK DQLVEPLGPGLKPPEERT EVAITFAKGLADKKISPK LVEKLTAYAMTIPFVRQG 21448 51816 C 21576 98 316  21449 51817 A 21577 35 247  21450 51818 A 21578 I 2391 LQLKMVACRAIGILSRFS RSRGYICRNFTGSSALLT YGVKGDVAADRINSPNS LSKELHSEFSEVMNEIWA RSAVLISSKPGCFIAGAI AACKVILQEVTQLSQEA EKP*KSPQSLIVGCPSNGS RTLRLPISSPQYRIATKVD *VPPEVFLGGPLPGRQG GCPKMGGVCLPAFGHEC SIR\ADRAKKMGLVD\QI GTQGLKPPEERTIEYLEE AKGLADKKISPKRDKGIU TA\YAMTIPFVRQQVYKI VRKQTKGLYPAPLKIIDV IEQGSDAGYLFEQKFG KESKALMGLYHGQVLCI GAPQKDVKHAILGAGL IAQVSVDKGLKTILKDAT DRGQQQVFKGINDKVKI TSPIERDSIFSNLTGQLDY KRPDMGELKAVFEDP*S VLKGKVGSR*FPDHWYS	
KSTKPIVAAINGSCLGGG SCQYRIATKDRKTVLGTI GALPGAGGTQRLPKMV LDMMLTGRSIRADRAKK DQLVEPLGPGLKPPEERT EVAITFAKGLADKKISPK LVEKLTAYAMTIPFVRQG  21448 51816 C 21576 98 316  21449 51817 A 21577 35 247  21450 51818 A 21578 I 2391 LQLKMVACRAIGILSRFS RSRGYICRNFTGSSALLT YGVKGDVAADRINSPNS LSKELHSEFSEVMNEIWA RSAVLISSKPGVCFIAGVAI AACK/TLQVEVTQLSQEVA EKP*KSPQSLIVGCPSNGS RTLRLPISSPQYRIATK\DS *VPPEVFLGGPLPGRQGG GCPKMGGVCLPAFGHEC SIRVADRAKVKMGLVDVQI GTQGLKPPEERTIEVLEE AKGLADKKISPKRDKG\I TAIYAMTIPFVRQQVYYK VRKQTKGLYPAPLKIIDV IEQGSDAGYLCESQKFGE KESKALMGLYHGQVLCE GAPQKDVKHLAILGAGL IAQVSVDKGLKTILKDAT DRGQQQVFKGLNDKVK TSF\ERDSIFSNLTGQLDY KRPDMGELKAVFEDPPS VLKGKVGSR*FPDHWYS	
SCQYRIATKDRKTVLGTI GALPGAGGTQRLPKMVC LDMMLTGRSIRADRAKK DQLVEPLGPGLKPPEERT EVAITFAKGLADKKISPK LVEKLTAYAMTIPFVRQC  21448 51816 C 21576 98 316  21449 51817 A 21577 35 247  21450 51818 A 21578 I 2391 LQLKMVACRAIGILSRFS RSRGYICRNFTGSSALLT YGVKGDVAADRINSPNS LSKELHSEFSEVMNEIWA RSAVLISSKPG\CFIAG\AI AACK\TLQ\EVTQLSQE\A EKP*KSPQSLIVGCPSNGG RTLRLPISSPQYRIATK\D *VVPE\VFLGGPLPGRQG GCPKMGGVCLPAFGHEC SIR\ADRAK\KMGLVD\QI GTQGLKPPEERTIEYLEE AKGLADKKISPKRDKGIL TA\YAMTIPFVRQ\VYK VRKQTKGLYPAPLKIIDV VREQTKGLYPAPLKIIDV IEQGSDAGYLCESQKFGF KESKALMGLYHGQVLCE GAPQKDVKHLAILGAGL IAQVSVDKGLKTILKDA1 DRGQQQVFKGLNDKVKI TSF\ERDSIFSNLTGQLDY KRPDMGELKAVF\EDPS VLKGKVGSR*FPDHWYS	
GALPGAGGTQRLPKMVQ LDMMLTGRSIRADRAKK DQLVEPLGPGLKPPEERT EVAITFAKGLADKKISPK LVEKLTAYAMTIPFVRQQ  21448 51816	
LDMMLTGRSIRADRAKK DQLVEPLGPGLKPPEERT EVAITFAKGLADKKISPK LVEKLTAYAMTIPFVRQU  21448 51816 C 21576 98 316  21449 51817 A 21577 35 247  21450 51818 A 21578 I 2391 LQLKMVACRAIGILSRFS RSRGYICRNFTGSSALLT YGVKGDVAADRINSPNS LSKELHSEFSEVMNEIWA RSAVLISSKPG\CFIAG\AI AACK\TLQ\EVTQLSQE\A EKP*KSPQSLIVGCPSNGG GCPKMGGVCLPAFGHEC SIR\ADRAK\KMGLVD\QI GTQGLKPPEERTIEYLEE AKGLADKKISPKRDKG\I TA\YAMTIPFVRQQVYKK VRKQTKGLYPAPLKIIDV IEQGSDAGYLCESQKFGE KESKALMGLYHGQVLCE GAPQKDVKHLAILGAGL IAQVSVDKGLKTILKDAT DRGQQQVFKGLNDKVKI TSF\ERDSIFSNLTGQLDY KRPDMGELKAVF\EDP*S VLKGKVGSR*FPDHWYS	
DQLVEPLGPGLKPPEERT EVAITFAKGLADKKISPK LVEKLTAYAMTIPFVRQUE 21448 51816 C 21576 98 316 21449 51817 A 21577 35 247 21450 51818 A 21578 I 2391 LQLKMVACRAIGILSRES RSRGYICRNFTGSSALLT YGVKGDVAADRINSPNS LSKELHSEFSEVMNEIWARSAVLISSKPGCFIAGIAI AACKITQIEVTQLSQEVA EKP*KSPQSLIVGCPSNGS RTLRIPISSPQYRIATKIDE SIRVADRAKKMGLVDVQI GTQGLKPPEERTIEVLEE AKGLADKKISPKRDKGIL TAVYAMTIPFVRQQVYKWARSAVLISKPKRDKGIL TAVYAMTIPFVRQQVYKWARSAVLISKPKRDKGIL TAVYAMTIPFVRQQVYKWARSAVLISKPKRDKGIL TAVYAMTIPFVRQQVYKWARSAVLIGAGLIAQVSVDKGLKTILKDAT DRGQQQVFKGLNDKVKITSFVERDSIFSNLTGQLDYKRPDMGELKAVFEQDPSVLKGKVGSR*FPDHWYS	
EVAITFAKGLADKKISPK LVEKLTAYAMTIPFVRQ0	
LVEKLTAYAMTIPFVRQQ	
21448   51816   C   21576   98   316	
21450 51818 A 21578 I 2391 LQLKMVACRAIGILSRFS RSRGYICRNFTGSSALLT. YGVKGDVAADRINSPNS LSKELHSEFSEVMNEIWA RSAVLISSKPG\CFIAG\AI AACK\TLQ\EVTQLSQE\A EKP*KSPQSLIVGCPSNGS RTLRLPISSPQYRIATK\DS *VPPE\VFLGGPLPGRQGG GCPKMGGVCLPAFGHEG SIR\ADRAK\KMGLVD\QI GTQGLKPPERTIEYLEE AKGLADKKISPKRDKG\I TA\YAMTIPFVRQVYKK VR\QTKGLYPAPLKIIDV IEQGSDAGYLCESQKFGE KESKALMGLVHGQVLCE GAPQKDVKHLAILGAGL IAQVSVDKGLKTILKDAT DRGQQQVFKGLNDKVKI TSF\ERDSIFSNLTGQLDY KRPDMGELKAVF\EDP*S VLKGKVGSR*FPDHWYS	
RSRGYICRNFTGSSALLT. YGVKGDVAADRINSPNS LSKELHSEFSEVMNEIWA RSAVLISSKPG\CFIAG\AI AACK\TLQ\EVTQLSQE\A EKP*KSPQSLIVGCPSNGS RTLRLPISSPQYRIATK\DS *VPPE\VFLGGPLPGRQGG GCPKMGGVCLPAFGHEG SIR\ADRAK\KMGLVD\QI GTQGLKPPEERTIEYLEE AKGLADKKISPKRDKG\I TA\YAMTIPFVRQQVYKK VRKQTKGLYPAPLKIIDV IEQGSDAGYLCESQKFGE KESKALMGLYHGQVLCE GAPQKDVKHLAILGAGL IAQVSVDKGLKTILKDAT DRGQQQVFKGLNDKVKI TSF\ERDSIFSNLTGQLDY KRPDMGELKAVF\EDP*S VLKGKVGSR*FPDHWYS	
YGVKGDVAADRINSPNS LSKELHSEFSEVMNEIWA RSAVLISSKPG\CFIAG\AI AACK\TLQ\EVTQLSQE\A EKP*KSPQSLIVGCPSNGS RTLRLPISSPQYRIATK\DS *VPPE\VFLGGPLPGRQGG GCPKMGGVCLPAFGHEG SIR\ADRAK\KMGLVD\QI GTQGLKPPEERTIEYLEE AKGLADKKISPKRDKG\I TA\YAMTIPFVRQQVYKK VRKQTKGLYPAPLKIIDV IEQGSDAGYLCESQKFGE KESKALMGLYHGQVLCE GAPQKDVKHLAILGAGL IAQVSVDKGLKTILKDAT DRGQQQVFKGLNDKVKI TSF\ERDSIFSNLTGQLDY KRPDMGELKAVF\EDP*S VLKGKVGSR*FPDHWYS	AFRIL
LSKELHSEFSEVMNEIWA RSAVLISSKPG\CFIAG\AI AACK\TLQ\EVTQLSQE\A EKP*KSPQSLIVGCPSNGS RTLRIPISSPQYRIATK\DS *VPPE\VFLGGPLPGRQG GCPKMGGVCLPAFGHEG SIR\ADRAK\KMGLVD\QI GTQGLKPPEERTIEYLEE AKGLADKKISPKRDKG\L TA\YAMTIPFVRQQVYKK VRKQTKGLYPAPLKIIDV IEQGSDAGYLCESQKFGE KESKALMGLYHGQVLCH GAPQKDVKHLAILGAGL IAQVSVDKGLKTILKDAT DRGQQQVFKGLNDKVKI TSF\ERDSIFSNLTGQLDY KRPDMGELKAVF\EDP*S VLKGKVGSR*FPDHWYS	NIHTS
RSAVLISSKPG\CFIAG\AI AACK\TLQ\EVTQLSQE\A EKP*KSPQSLIVGCPSNGS RTLRLPISSPQYRIATK\DS *VPPE\VFLGGPLPGRQGG GCPKMGGVCLPAFGHEG SIR\ADRAK\KMGLVD\QI GTQGLKPPERTIEYLEE AKGLADKKISPKRDKG\L TA\YAMTIPFVRQQVYKK VRKQTKGLYPAPLKIIDV IEQGSDAGYLCESQKFGE KESKALMGLYHGQVLCE GAPQKDVKHLAILGAGL IAQVSVDKGLKTILKDAT DRGQQQVFKGLNDKVK TSF\ERDSIFSNLTGQLDY KRPDMGELKAVF\EDP*S VLKGKVGSR*FPDHWYS	<\VNT
AACK\TLQ\EVTQLSQE\A EKP*KSPQSLIVGCPSNGS RTLRLPISSPQYRIATK\DS *VPPE\VFLGGPLPGRQGG GCPKMGGVCLPAFGHEG SIR\ADRAK\KMGLVD\QI GTQGLKPPEERTIEYLEE AKGLADKKISPKRDKG\L TA\YAMTIPFVRQQVYKK VRKQTKGLYPAPLKIIDV IEQGSDAGYLCESQKFGE KESKALMGLYHGQVLCE GAPQKDVKHLAILGAGL IAQVSVDKGLKTILKDAT DRGQQQVFKGLNDKVK TSF\ERDSIFSNLTGQLDY KRPDMGELKAVF\EDP*S VLKGKVGSR*FPDHWYS	SDQI
EKP*KSPQSLIVGCPSNGS RTLRLPISSPQYRIATK\DS *VPPE\VFLGGPLPGRQGG GCPKMGGVCLPAFGHEG SIR\ADRAK\KMGLVD\QI GTQGLKPPERTIEYLEE AKGLADKKISPKRDKG\L TA\YAMTIPFVRQQVYKK VRKQTKGLYPAPLKIIDV IEQGSDAGYLCESQKFGE KESKALMGLYHGQVLCE GAPQKDVKHLAILGAGL IAQVSVDKGLKTILKDAT DRGQQQVFKGLNDKVKI TSF\ERDSIFSNLTGQLDY KRPDMGELKAVF\EDP*S VLKGKVGSR*FPDHWYS	ITML
RTLRLPISSPQYRIATK\DO *VPPE\VFLGGPLPGRQGG VPKMGGVCLPAFGHEG SIR\ADRAK\KMGLVD\QI GTQGLKPPEERTIEYLEE AKGLADKKISPKRDKG\L TA\YAMTIPFVRQQVYKK VRKQTKGLYPAPLKIIDV IEQGSDAGYLCESQKFGE KESKALMGLYHGQVLCE GAPQKDVKHLAILGAGL IAQVSVDKGLKTILKDAT DRGQQQVFKGLNDKVKK TSF\ERDSIFSNLTGQLDY KRPDMGELKAVF\EDP*S VLKGKVGSR*FPDHWYS	QR\IV
*VPPE\VFLGGPLPGRQGG GCPKMGGVCLPAFGHEC SIR\ADRAK\KMGLVD\QI GTQGLKPPEERTIEYLEE AKGLADKKISPKRDKG\L TA\YAMTIPFVRQQVYKK VRKQTKGLYPAPLKIIDV IEQGSDAGYLCESQKFGE KESKALMGLYHGQVLCE GAPQKDVKHLAILGAGL IAQVSVDKGLKTILKDAT DRGQQQVFKGLNDKVKI TSF\ERDSIFSNLTGQLDY KRPDMGELKAVF\EDP*S VLKGKVGSR*FPDHWYS	CLGR
GCPKMGGVCLPAFGHEC SIR\ADRAK\KMGLVD\QI GTQGLKPPEERTIEYLEE AKGLADKKISPKRDKG\L TA\YAMTIPFVRQQVYKK VRKQTKGLYPAPLKIIDV IEQGSDAGYLCESQKFGE KESKALMGLYHGQVLCE GAPQKDVKHLAILGAGL IAQVSVDKGLKTILKDAT DRGQQQVFKGLNDKVKI TSF\ERDSIFSNLTGQLDY KRPDMGELKAVF\EDP*S VLKGKVGSR*FPDHWYS	KTVI
SIR\ADRAK\KMGLVD\QI GTQGLKPPEERTIEYLEE AKGLADKKISPKRDKG\L TA\YAMTIPFVRQQVYKK VRKQTKGLYPAPLKIIDV IEQGSDAGYLCESQKFGE KESKALMGLYHGQVLCE GAPQKDVKHLAILGAGL IAQVSVDKGLKTILKDAT DRGQQQVFKGLNDKVK TSF\ERDSIFSNLTGQLDY KRPDMGELKAVF\EDP*S . VLKGKVGSR*FPDHWYS	HTK
GTQGLKPPEERTIEYLEE AKGLADKKISPKRDKG\L TA\YAMTIPFVRQQVYKK VRKQTKGLYPAPLKIIDV IEQGSDAGYLCESQKFGE KESKALMGLYHGQVLCE GAPQKDVKHLAILGAGL IAQVSVDKGLKTILKDAT DRGQQQVFKGLNDKVKI TSF\ERDSIFSNLTGQLDY KRPDMGELKAVF\EDP*S . VLKGKVGSR*FPDHWYS	LTGR
AKGLADKKISPKRDKG\L TA\YAMTIPFVRQQVYKK VRKQTKGLYPAPLKIIDV IEQGSDAGYLCESQKFGE KESKALMGLYHGQVLCE GAPQKDVKHLAILGAGL IAQVSVDKGLKTILKDAT DRGQQQVFKGLNDKVKI TSF\ERDSIFSNLTGQLDY KRPDMGELKAVF\EDP*S	VGTP
TA\YAMTIPFVRQQVYKK VRKQTKGLYPAPLKIIDV IEQGSDAGYLCESQKFGE KESKALMGLYHGQVLCE GAPQKDVKHLAILGAGL IAQVSVDKGLKTILKDAT DRGQQQVFKGLNDKVKI TSF\ERDSIFSNLTGQLDY KRPDMGELKAVF\EDP*S . VLKGKVGSR*FPDHWYS	/AI\TF
VRKQTKGLYPAPLKIIDV IEQGSDAGYLCESQKFGE KESKALMGLYHGQVLCE GAPQKDVKHLAILGAGL IAQVSVDKGLKTILKDAT DRGQQQVFKGLNDKVKI TSF\ERDSIFSNLTGQLDY KRPDMGELKAVF\EDP*S . VLKGKVGSR*FPDHWYS	VEKL
IEQGSDAGYLCESQKFGE KESKALMGLYHGQVLCE GAPQKDVKHLAILGAGL IAQVSVDKGLKTILKDAT DRGQQQVFKGLNDKVKI TSF\ERDSIFSNLTGQLDY KRPDMGELKAVF\EDP*S . VLKGKVGSR*FPDHWYS	VEEK
KESKALMGLYHGQVLCH GAPQKDVKHLAILGAGL IAQVSVDKGLKTILKDAT DRGQQQVFKGLNDKVKI TSF\ERDSIFSNLTGQLDY KRPDMGELKAVF\EDP*S . VLKGKVGSR*FPDHWYS	VKTG
GAPQKDVKHLAILGAGL IAQVSVDKGLKTILKDAT DRGQQQVFKGLNDKVKI TSF\ERDSIFSNLTGQLDY KRPDMGELKAVF\EDP*S . VLKGKVGSR*FPDHWYS	LVMT
IAQVSVDKGLKTILKDAT DRGQQQVFKGLNDKVKI TSF\ERDSIFSNLTGQLDY KRPDMGELKAVF\EDP*S . VLKGKVGSR*FPDHWYS	KNKF
DRGQQQVFKGLNDKVKI TSF\ERDSIFSNLTGQLDY KRPDMGELKAVF\EDP*S . VLKGKVGSR*FPDHWYS	MGAG
TSF\ERDSIFSNLTGQLDY KRPDMGELKAVF\EDP*S VLKGKVGSR*FPDHWYS	LTAL
KRPDMGELKAVF\EDP*S . VLKGKVGSR*FPDHWYS	KAL
VLKGKVGSR*FPDHWYS	QGF*
	LSHR
1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	LPSNT
SALPISEIAAVQQKT*EG*	LGMH
YF\SPVDKMQLL\EIITTE	TSKD
T\SASAVA\VGLKQGK\VI	(V\VK
DG\PGLLVTYQGVFAPMI	1S\EVI
	rtsfg
F\PVG\AATLV\DEVGCGN	(LAK
HV\AEDLGKVFG\ERFGG	<b>GNPE</b>
L\LTQMVSKGFLG\RKSG	GLLT
SSGRVVKRKILNSTMD\S	*ASL
21451 51819 A 21579 119 387 VEDSKEIFSSA*DYLSTHS	KHVP
ASHESYHVNPNLCNPVA	TSGA
HSIV*KWPSWLGTVAHSO	CNPSN
FGRPRRADCLSSRSSRPA	WPTR
21452 51820 B 21580 73 3177	

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)  SGEPERALRGAAMRPSGTAGA
21453	51821	A	21581	1399	3140	ALLALLAALCPASRALEEKKVC QGTSNKLT\QLGTFE\DHFPQPS QRMF\NNCEVV\LGNLEITLCER NYDLSFLKTI\QEVAGYVLIALN TVERIPLENLQIIRGNMYYENSY ALAV\LFNYDANKTGLKELPMR NLQEILHGAVRFSNNPALCNVE SIQWRDIVSSDFLSNMSMDFQN HLGSCQKCDPSCPNGS\CWGAG EENCQKLTKIICAQQCSGRCRG KSPSDCCHNQ
21454	51822	В	21582	82	305	
21455	51823	Α	21583	122	219	YAKRIYDVYYVHTAYTCTFHA *KIRGMTQ*NW
21456	51824	A	21584	404	1228	GLAAFLARARGRRRKELPGSPR PLPQLPAGGPSPRWRAMRPSQR RPPTLPMAASPAHWPRESLVLY HWTQSFSSQKVRLVIAEKGLVC EERDVSLPQSEHN/EMPWFMRL NLGEEVPV\IIHRDNIIQ/CTMDQ IHLTMWERTFHRERHVGALMP EVGSRAA\PRVLQY\RELLDALP MDAYTHGCILHPELTTDSMIPK YATAEIRRHLANATTDLMKLD HEEEPQLSEPYLSKQKKLMAKI L\EHDDVSYLKKILGELAMVLD QIEGGAGRKRKLENE
21457	51825	A	21585	3	1194	KNLSVKLSENGTLKNCVVADS ETDIDVYMYTRVYVYIPRIENS WNDTIKLVTGATISAVGGDIDA QGEWAFPLLGIPSLSPGPEWKV RVVIPDKGLVCEERDVSLPQSE PKEPWFMRINLGEEVPVIIHRD NIISDYDQIIDYVERTFTGEHVV ALMPEVGSLQHARVLQYRELL DALPMDAYTHGCILHPELTTDS MIPKYATAEIR/SDI*PNATTDL MKLDHEEPQLSEPYLSKQKKL MAKILEHDDVSYLKKILGELA MVLDQIEAELEKRKLENEGQK CELWLCGCAFTLADV\SLGATL HRLKFLGLSKKYWEDGSRP\NL QSFFERVQRRFAFRKVLGDIHT TLLSAVIPN\AFRLVKRKPPSFFR ASFLMGSLG\GMGYFALHGTSR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21458	51826	Α	21586	1	929	MDPSIYFLPSKREGHLQAYIRDS
						VPEDPLILIFFSASSDQRPWLKA
						PHLPALRGFRGGGAGGAQSR
						TTEDYVAWAKGNSPTSPSPRSR
		ł				GGCSAKLSGPRRPPRRPGRLFN
İ						MAASANSVSRAWRTESPEGLE
					}	SLHSTSLSAISVSRPEGGAELSD
1						TADTMGFGDLKSPAGLQV\PTI
						TWRTRATSRGMCHHKQMWQY
						LKPCPAHRLPTCVMPYV/GYNH
						IKSYEKEKASLPGVKKALGKYG
						PADVEDTTGSGATDSKDDDDID
		-				LFGSDDEEESEEAKRLRE/DTSC
						TI*IKESQKTCTCCQVFHLTRCE
21459	51827	Α	21587	1	2405	MALRVVRSVRAVLCSLHVACE
						KVGMQIPRFCYHERLSVAGNC
						RMCLVEIEKAPKDQSMMFGND
						RSRFLEGKRAVEDKNIGPLVKT
						IMTRCIQCTRCIRFASEIAGVDD
						LGTTGRGNDMQVGTYIEKMFM
						SELSGNIIDICPVGALTSKPYAFT
						ARPWETRKTESIDVMDAVGSNI
						VVSTRTGEVMRILPRMHEDINE
						EWISDKTRFAYDGLKRQRLTEP
						MVRNEKGLLTYTSWEDALSRV
						AGMPPALALQSAEIAASARPPP
						RLGSEEPLCLAAHRLGCEEPLC
					}	LAASLESEERLCPAAIPPGNLRY
						VFISSAKTDKYSKLAPVEWVTV
						EKVPENVEVTLELGNRQSRDR
						VSRIPGNAIVIWFTGFKWKKTV
						TTLWFLNLAIADFIFLLFLPLYIS
	ļ					YVAMNFHWPFGIWLCKANSFT
						AQLNMFASVFFLT/GDQPGPLY
						PLDPSCLISSASNPQELSDCHYI
						HLAFGFSNWRSCPVLPGHCGV
						Q*SYSLL*QFSEA*S*PHFDQAP
						CSDLGEIYHWLSLPFANNEYLL
1		1				LVSHLQGEEAKHPDLQ*AFLDN
1						SGCGCGLCGLLDSLSPV*HLGA
						HHSPQ*LFPPCDAGWNPPLHWF
1						GIPQ*LLEPHPLCPN**EVPSSLP
						VLSC*DTQVHTVGSQLFW\QVL
						KEAKKPMVVLGSSALQRNDGA
						AILAAVSSIAQKIRMTSGVTGD
		<u> </u>		L	<u> </u>	WKVMNILHRIASQVAALDLGY

2750

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
21460	51828	A	21588	1	2065	MLRIPVRKALVGLSKSPKGCVR
						TTATAASNL\IRVFVDGQSLM\V
						EPGTTVLQACE\RIGMQIPRFCY
						HERLSVAGNCRMCLVEIEKAPK
						DQSMMFGNDRSRFLEGKRAVE
		1				DKNIGPLVKTIMTRCIQCTRCIR
						FASEIAGVDDLGTTGRGNDMQ
		l				VGTYIEKMFMSELSGNIIDICPV
						GALTSKPYAFTARPWETRKTES
			[			IDVMDAVGSNIVVSTRTGEVM
						RILPRMHEDINEEWISDKTRFA
		i				YDGLKRQRLTEPMVRNEKGLL
ļ						TYTSWEDALSRVAGMLQSFQG
						KDVAAIAGGLVDAEALVALKD
						LLNRVDSDTLCTEEVFP\TAGA
					j	GTDLRSNYLLNTTIAGV\EEAD
						VVLLVGTNPRF\EAPLF*C*EFR
[						KSW\LHND\LKVALIRAVPVDL\
•						TYTYDHLG\DSPKILQ\DIASGK/
						SHPFSQVLKEAKKPMVV\LGSS
						ALQRNDGAAILAAVSSIAQKI\R
						MTSGVTGDWKVMNILHRIASQ
						VAVL\DLGYKPGVEAIRKNPPK
						VLFLLGADGGCITRQDLPKDCF
						IIY\QGHHGDVGAPIADVILPGA
						AYTEKSATYVNTEGRAQQTKV
						AVTPPGLAREDWKIIRALSEIAG
						MTLPYDTLDQVRNRLEEVSPNL
		ł				VRYDDIEGANYFQQANELSKL
}		1				VNQQLLADPLVPPQLTIKDFYM
						TGEVTEINEALAENPGLVNKSR
						YEDGWLIKMTLSNPS

2751

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21461	51829	Α	21589	56	2352	ATRPPRGSSWCRQFSRTASAAP
ŀ						GRSNMLRIPVRRALVGLSKSPK
				:		GCVRTTATAASNLIEVFVDGQS
į						VMVEPGTTVLQACEKVGMQIP
						RFCYHERLSVAGNCRMCLVEIE
						KAPKVVAACAMPVMKGWNIL
	i	ĺ		:		TNSEKSKKAREGVMEFLLANH
İ						PLDCPICDQGGECDLQDQSMM
						FGNDRSRFLEGKRAVEDKNIGP
						LVKTIMTRCIQCTRCIRFASEIA
						GVDDLGTTGRGNDMQVGTYIE
						KMFMSELSGNIIDICPVGAL\TS
						KPYAFTARPWETRKTESIDVMD
						AVGSNIVVSTRTGEVMRILPRM
						HEDINEEWISDKTRFAYDGLKR
						QRLTEPMVRNEKGLLTYTSWE
		ļ				DALSRVAGMLQSFQGKDVAAI
					:	AGGLVDAEALVALKDLLNRVD
						SDTLCTEEVFPTAGAGTDLRSN
						YLLNTTIAGVEEADVVLLVGTN
						PRFEAPLFNAWIRKSWLHNDLK
						VALIGSPVDLTYTYDHLGDSPK
		ĺ				ILQDIASGSHPFSQVLKEAKKP
						MVVLGSSALQRNDGAAILAAV
1						SSIAQKIRMTSGVTGDWKVMNI
						LHRIASQVAALDLGYKPGVEAI
						RKNPPKVLFLLGADGGCITRQ\
ł						DLPKDCFIIYQGHHGDVGAP\IA
						DVILPGAAYTN\KSATYVNTEG
						*ASASIMLAVTT\PGLSREDW\KI
						IRALSEIAGMT/LFPYDTL/DIQV
						RNQIGKKSSPNLV\RYDDIEG\A
21462	51830	Α	21590	246	761	MRLQGAIFVLLPHLGPILVWLF
						TRDHMSGWCEGPRMLSW\CPF
						LQSLICLLQTAI\YSVVG*VLVS
						HTWPWYPMGWEQVLYI*ITYA
				1		VQLTISWTVLVLFFTVHNPGLV
						RHTVLSSRSNVGEGETTWLQK
						HILAALLLLPYLAWLTVTSALT
						YHLWRDSLCPVHQPQPTEKSD
21463	51831	Α	21591	1	1497	
21464	51832	Α	21592	<u> </u>	1731	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	1			sequence		
21465	51833	A	21593	1	6626	PRSLCFSLWAEAAVLADGGLR
		İ				RRRRLLRGTMSASFVPNGASLE
						DCHCNLFCLADLTGIKW*KYV
į		•				WGGPTSAPILFPVTEEHPILSSSS
						RCLTAHVLGVRRRDQRPERRE\
						L*IFWWGEDP\VLLTLFTMTYQ
		ŀ				KKKMECGRMDFPMNAVLCFS
						KAVHNLLERCLMNRNFVRIGK
						WFVKPYEKDEKPINKSEHLSCS
						FTFFLHGDSNVCTSVEINQHQP
						VYLLSEEHITLAQQSNSPFQVIL
						CPFGLNGTLTGQAFK
21466	51834	Α	21594	1263	1906	AVRGMVRNARAVLKTGGTFLS
			,			PALEPLVAGGSGG/EFSQLGAEE
						RKAIARF/AIDHVDRRVPVLIGT
		ŀ				GG/TNARETIELSQHAQQAGRT
•						ASTIDSVAHLRSMIHTVK/GAHP
						HFTVLCGYDDHLF/NTLLLGGD
						GAISASGNFAPQVSVNLLKAW
	}					RDGDVAKAAGYHQTASIPDQLI
						HPYESRLFKCQHVLTRGITSF*L
	ļ	<u> </u>				LCPPGSLRRQGISTYHSLLHN
21467	51835	В	21595	113	484	VOCTA IETIDAN A DOCKINIA DO
21468	51836	A	21596	300	780	KSGTNETIRRKARGGKIK\MDD
						LRG\PRCQ*EPWKRLLMTIMPS CLHLWAQNTTVSILSFVDKDLL
						EPGCSVLLNHKVHAVIGVLMD
i	1					DTDPLVTVMKVEKAPQETYAD
						IGGLDNQIQEIKESVELPLTHPE
						YYEEMGIKPPKGVHPFGPTWPR
						VKPFCPRQ
21469	51837	A	21597	2	840	AIEHCQSGDNPESRRGFLLQQW
21105	15.05.	` `		-		LGRNPALVPHPGRTGHSQPPVT
						FHRHPSDCQRSPAGRFKGGPSH
						RGOPPPFHKSPPMTTASHLARC
	}					ARTYHQ\DSEAAINR\QINLE\LY
						ASYVY\LSMSYYFDRD\DVALK
						NF\AKYFLHQSHEER\EHC*ENL
						MKACRTNEGWPNLSFQDIKET
						KTCDD\WESGLNA\ME\CALHFE
						KKC*IKSLLEL\HKL\ATDKN*PP
						ICVDFI*DTFTLNEQ/V*KAIKRI
						WGDH/V*PKLWRKMGSAPNLG
]						FGEYLF*QSTPWGDSDNES
21470	51838	A	21598	3	390	
					·	

2753

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21471	51839	A	21599	11	483	FQLEVTPQNITLNPGFGGPVFS
21471	31037	``	21377	,	103	WRDQAVLRQDGVVVTINKKR
						NLVVSVDDGGTFEVVLHRVWK
						GSSVHQDFLGFYVLDSHRMSA
						RTHGLLGQFFHPIGFEVSDIHPG
						SDPTKPDATMVG\LQKDYSKDP
						WHGAEVSCWFIHNNGAGLIDG
						AYTDYIVPDIF
21472	51840	A	21600	1	1521	
21473	51841	Α	21601	1	837	
21474	51842	Α	21602	3	310	EVRANLSSQALRMWLDYGFVT
						PLTSMSIRGMADQDGLKPTIDK
						PSEDSPPLEMLGPRRTFVLSALQ
						PSPTHSSSNTQRLPDRVTGVDT*
						VVT*PYVLCWNLC
21475	51843	Α	21603	62	462	MPLPLPSAFVLSALQPSPTHSSS
						NTQRLPDRVTGGFSVNGQLIGN
						KARSPGQ\HDGTYFGRLGIANP
						ATDFQLEVTPQNITLNPGFGGP
						VFSW\RDQAVLRQDGVVVTI\N
			1			KKRNLVVSVDDGGTFEVVLHR
						VWK
21476	51844	Α	21604	37	2883	SMDGAMGPRGLLLCMYLVSLL
			•			ILQAMPALGSATGRSKSSEKRQ
						AVDTAVDGVFIRSLKVNCKVTS
ŀ						RFAHYVVTSQVVNTANEAREV
		1				AFDLEIPKTAFISDFAVTADGNA
						FIGDIKDKVTAWKQYRKAAIS*
						EKA\GLVRASGRTMEQFTIHLT
						VNPQSKVTFQLTYEEMLKRNH
		1				MQYEIVIKVKPKQLVHHFEIDV
						DIFEPQGISKLDAQASFLPKELA
						AQTIKKSFSGKKTSNGSHSRRF
		_				QGHVLFRPTVSQQQ
21477	51845	A	21605	1	1008 435	
21478 21479	51846 51847	A	21606 21607	1	676	IRHQGAAAQEGDAETPGSVERR
214/9	31047	^	21007		070	GRRAGAEDGMSQAPGAQPSPP
				1		TVYHERQRLELCAVHALNNVL
						QPAA\FSQEAADEICKRLAPDSR
ł						LNPHRSLLGTGNYDVNVIMAA
		1				LOGLGLAAVWWDRRRPLVPS
					:	WPCPKGLG*ILN\LPSPVSLGLL
						SLPLRRRHLRWPCARL/VTVSY
1		1				YNLDSK\LRAPEGPGGLRTE/LR
						GLSLAAALAQGLCEVLLVVTK
			1			EVEEKGSWLRTD
21480	51848	A	21608	89	316	ELLSPLLVAPGPSDLTVTGLKSF
21400	31040	^	21008	0,7	310	SSPSLCSLNTLLAGTSRLMVA*L
						YESLGQGEDLGLLTPGPEFLSIS
		1				SYCNFM
L	J		<u>L </u>	<u> </u>	L	010141141

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21481	51849	В	21609	1	1920	
21482	51850	A	21610	1	1727	LPRVLGHSGHRGGLSSRRSQGL
						VSRFRLRALCGAGSDMWRLPG
						LLGRALP\RTLGP\SLWRVTPKS
						SSPDGPQTTSST\LLVPVPNLDR
		-				SGPHGPGTSGGPRSHGWKDAF
						QWMSSRVSPNTLW\DAISWGTL
						AVLALQLGR\HIHFQAFLAAGP\
						KRVE\HCS\WHSPL\DRFFSSP\L
						WHPCSSLRQHILPSPDGPAPRHT
						GLK/ENPRLGKE\EASAQPRNFS
1						1
						HNSLRGARP\QDPSEEG\PGDFG
						FLHASSSIESEAKPAQPQPTGEK
						EQDKSKTLSLEEAVTSIQQLFQL
						SVSIAFNFLGNQMDPAPRQEGT
		1				ENMKSGDHTAAFSYFQKAAAR
				1		GYSKAQYNAGLCHEHGRGTPR
		ł				DISK/SV*KAGLYYQLAASQGH
						SLAQYRYARCLLRDPASSWNPE
		ŀ				RQRAVSLLKQAADSGLREAQA
						FLGVLFTKEPY\LDEQRAVKYL
						WLAANNGDSQSRYHLGICYEK
		<b> </b>				GLGVQRNLGEALRCYQQSAAL
						GNEAAQERLRALFSMGA\ASPG
						APTNLTVTGLKSFSSPSLCSLNT
		1				LLAGTSR\LPH\ASSTGNLGLLC
						RSGHLGASLEASSRAIPPHPYP\
						LEKN\VVRLGFG
21483	51851	Α	21611	1104	1680	LATPDVRYCYLSTYTSLPNPAR
						RFFLFSLCCKFSHSH\KTATVKK
		ŀ				YNKSEYPWDLVKAHLQGAFTS
						NITFDISELQNKILDLNRHTQEF
}						QPSLEDWTEFQEGLESLNPWTY
						LKYHSNILYVVLGVMLFCLCLL
		1				FIVCKIGWTANRRMRAAQPGL
						TFFQLIHKQGDMQGAEGPWDV
						TNSAFHWRLYDQTANCLS
21484	51852	A	21612	364	824	LCIILFKQLSTWRTGCAQPTKR
}						GYTELQTEGPMLKSMFLWLGE
1		1				QVQILQL\HSNCIVILTTLV/FCV
ŀ						TNLE*FIVCKIGWTANQKMGA
				1	'	AKPGLTFFLLICK\QKGDM*GA
1		1				KGPWDVINSAFH*RM
21485	51853	c	21613	196	504	THE WEST THE THE
21486	51854	A	21614	3	142	
21700	121027	112	[= 1 U 1 T	<u>  -                                   </u>	1	l

SEQ ID NO:	SEQ ID NO: of peptide	Met	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
NO.	sequence	1,00	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	}			sequence		· i
21407	51055		01615	1170	2295	IVINA METCOCORNAL CALLED
21487	51855	Α	21615	1178	2285	IVWLMTTGGQCRY*LSALIKL*
						DR/WFQRLCIHVIQRLRPVHAH
	1					LYLQPGMEDGNSFRQMFRSTR
						SLHIPTRDLPLSPDTTVVLHQVY
						NVLLGLLSRAKLYVDAAVHGT
	İ	1				TKLVPYFSFMTYCLISKTEKLM
1						FSTYFMDLWNLFQPKLSEPAIA
		1				TNHNKQALLSFWYNVCADCPE
						NIRLIVQNPVVTKNIAFNYILAD
						HDDQDVVLFNRGMLPAYYGIL
					·	RLCCEQSPAFTRQLASHQNIQW
						AFKNLTPHASQYPGAVEELFNL MQLFIAQRPDMREEELEDIKQF
						KKTTISCYLRCLDGRSCWTTLIS
						AFRILLESDEDRLLVVFNRGLIL
ļ				į		MTEGKD\TFMIVCC*TTSFLIISS
						l
21400	51056		21616	1	436	SIYYAELQSTVKNLLKH
21488 21489	51856 51857	A	21616	369	508	
21490	51858	A	21618	2150	3120	DLRALALLLSVHTPKQLNPALI
21490	131030	<u>Γ</u>	21016	2130	3120	PTLQELLSKCRTCLQQRNSLQE
						QEAKERKTKALALWTTIITFRV
		1				GGGSNTLGVTGLRVVCSAEPPK
						YKC*KQN*LPTSPPNVILMTFRE
		l			1	VSLLACVFTDDEGATPIKRRRV
						SSDEEHTVDSCISDMKTETREV
						LTPTSTSDNETRDSSIIDPGTEQ
		l				DLPSPENSSVKEYRMEVPSSFSE
						DMSNIRSQHAEEQSNNGRYDD
						CKEFKDLH\CSKDSTLA\EEESE\
						FPSTSISAVLSDLADLRSCDGQA
						LPSQDPEVALSLSCGHSRGLFS
		l				HMQQHDILDTLCRTIESTIHVVT
		1				RISGKGNQAAS
21491	51859	A	21619	2	4049	EEENGREYKFDVSSELLEWIWH
						DNMQFLQDKNIFEHTYFGFMW
		1				QLCSCIPSTLPDPKAVSLMTAK
		l				LSTSFVLETFIHSKEKPTMLQWI
						ELLTKQFNNSQAACEWFLDRM
						ADDDWWPMQILIK\CPNQIVRQ
						MFQRLCIHVIQRLRPVHAHLYL
1		}				QPGMEDGSDDMDTSVEDIGGR
						SCVTRFVRTLLLIMEHGVKPHS
1						KHLTEYFAFLYEFAKMGEEESQ
						FLLSLQAISTMVHFYMGTKGPE
1						NPQVEVLSEEEGEEE
21492	51860	Α	21620	58	219	
			<u> </u>		L	<u></u>

NO: of peptide sequence hod in USSN 09/540,217 codon for peptide sequence codon for peptide sequence codon for last amino acid of peptide sequence deletion, \=possible nucleon for peptide sequence codon for last amino acid of peptide sequence deletion, \=possible nucleon for peptide sequence codon for last amino acid of peptide sequence codon for last amino acid of peptide sequence codon for last amino acid of peptide sequence codon for last amino acid of peptide sequence codon for last amino acid of peptide sequence codon for last amino acid codon for last amino acid codon for last amino acid of peptide sequence codon for last amino acid codon for last amino	ASRSRENEGL PASSVQAWEA PKGVEPPAEE PTSRKDPKFE ATLLPEHIVVR HHLFRKWAAL SQKHKLTVVL GRPSGA/WEST VALAKRFQP
21493 51861 A 21621 195 1187 RVPRTWLLPSFC/ SEFGASQTVTPSP MEPEFLYDLLQL ELSKGVCKKYLP ELQKVQQEWINA SLEEDMFDGLILF KLEAEDIALTATS EAVNRS\CSWRSG WNKDLLSTLHLL DLSLP/D/HTNVQ NFVNQKLDRLGL GTGVILLLLIGQL YLTPNSPAEMVS	ASRSRENEGL PASSVQAWEA PKGVEPPAEE PTSRKDPKFE ATLLPEHIVVR HHLFRKWAAL SQKHKLTVVL GRPSGA/WEST VALAKRFQP VEVITIEVAA
21493 51861 A 21621 195 1187 RVPRTWLLPSFC/SEFGASQTVTPSPMEPEFLYDLLQLELSKGVCKKYLPELQKVQQEWINASLEEDMFDGLILFKLEAEDIALTATSEAVNRS\CSWRSGWNKDLLSTLHLLDLSLP/D/HTNVQNFVNQKLDRLGLGTGVILLLLIGQLYLTPNSPAEMVS	PASSVQAWEA PKGVEPPAEE PTSRKDPKFE ATLLPEHIVVR HHLFRKWAAL SQKHKLTVVL GRPSGA/WEST VALAKRFQP VEVITIEVAA
SEFGASQTVTPSP MEPEFLYDLLQL ELSKGVCKKYLP ELQKVQQEWINA SLEEDMFDGLILH KLEAEDIALTATS EAVNRS\CSWRSG WNKDLLSTLHLL DLSLP/D/HTNVQ NFVNQKLDRLGL GTGVILLLLIGQL YLTPNSPAEMVS	PASSVQAWEA PKGVEPPAEE PTSRKDPKFE ATLLPEHIVVR HHLFRKWAAL SQKHKLTVVL GRPSGA/WEST VALAKRFQP VEVITIEVAA
MEPEFLYDLLQL ELSKGVCKKYLP ELQKVQQEWINA SLEEDMFDGLILH KLEAEDIALTATS EAVNRS\CSWRSG WNKDLLSTLHLL DLSLP/D/HTNVQ NFVNQKLDRLGL GTGVILLLLIGQL YLTPNSPAEMVS	PKGVEPPAEE PTSRKDPKFE ATLLPEHIVVR HHLFRKWAAL SQKHKLTVVL GRPSGA/WEST VALAKRFQP VEVITIEVAA
MEPEFLYDLLQL ELSKGVCKKYLP ELQKVQQEWINA SLEEDMFDGLILH KLEAEDIALTATS EAVNRS\CSWRSG WNKDLLSTLHLL DLSLP/D/HTNVQ NFVNQKLDRLGL GTGVILLLLIGQL YLTPNSPAEMVS	PKGVEPPAEE PTSRKDPKFE ATLLPEHIVVR HHLFRKWAAL SQKHKLTVVL GRPSGA/WEST VALAKRFQP VEVITIEVAA
ELSKGVCKKYLP ELQKVQQEWINA SLEEDMFDGLILH KLEAEDIALTATS EAVNRS\CSWRSG WNKDLLSTLHLL DLSLP/D/HTNVQ NFVNQKLDRLGL GTGVILLLLIGQL YLTPNSPAEMVS	PTSRKDPKFE ATLLPEHIVVR HHLFRKWAAL SQKHKLTVVL GRPSGA/WEST VALAKRFQP VEVITIEVAA
ELQKVQQEWINA SLEEDMFDGLILH KLEAEDIALTATS EAVNRS\CSWRSC WNKDLLSTLHLL DLSLP/D/HTNVQ NFVNQKLDRLGL GTGVILLLLIGQL YLTPNSPAEMVS	ATLLPEHIVVR HHLFRKWAAL SQKHKLTVVL GRPSGA/WEST VALAKRFQP VEVITIEVAA
SLEEDMFDGLILH KLEAEDIALTATS EAVNRS\CSWRSG WNKDLLSTLHLL DLSLP/D/HTNVQ NFVNQKLDRLGL GTGVILLLLIGQL YLTPNSPAEMVS	HHLFRKWAAL SQKHKLTVVL GRPSGA/WEST VALAKRFQP VEVITIEVAA
KLEAEDIALTATS EAVNRS\CSWRSG WNKDLLSTLHLL DLSLP/D/HTNVQ NFVNQKLDRLGL GTGVILLLLIGQL YLTPNSPAEMVS	SQKHKLTVVL GRPSGA/WEST VALAKRFQP VEVITIEVAA
EAVNRS\CSWRSO WNKDLLSTLHLL DLSLP/D/HTNVQ NFVNQKLDRLGL GTGVILLLLIGQL YLTPNSPAEMVS	GRPSGA/WEST VALAKRFQP VEVITIEVAA
WNKDLLSTLHLL DLSLP/D/HTNVQ NFVNQKLDRLGL GTGVILLLLIGQL YLTPNSPAEMVS	VALAKRFQP VEVITIEVAA
DLSLP/D/HTNVQ NFVNQKLDRLGL GTGVILLLIGQL YLTPNSPAEMVS:	VEVITIEVAA
NFVNQKLDRLGL GTGVILLLIGQL YLTPNSPAEMVS	
GTGVILLLIGQL YLTPNSPAEMVS	SMONI INTOW I
YLTPNSPAEMVS	, ,
GRGPAQLPC/LAI	
KSTLRVLYGLFC	KHIQKAHRD
RTPHGAPN	DDDDTDNILD
21494 51862 A 21622 96 1350 AAGLLPPGLVPEI	
FGIQGPPFALSRP	i
WEAMEPEFLYDL	`
AEEELSKGGKKK	i
KFEELQKPA\VLN	
EHIVVRSLEEDMI	i i
QRLAALKLEAED	
HKLHSWCWRP*7	
PSGA/WESIFNKD	
LAKRFQPDLSLPT	•
STKSGLKSEKLER	EQLTEYSTDK
DEPPKDVFDELFI	1
VKEAIVNFVNQK	- 1
	LLLIGQLEGFF
LHLKEFYLTPNSF	PAEMLHNVTL
ALELL/IGRGPAQ	LPC/LALK/Ti
VNKDAKSTLRVL	.YGLFCKHTH
KAHRDRTPPWSP	ELTLTASKAQ
SLPVSPAGGPEAA	AGCPPTVPLFP
21495 51863 A 21623 1 87	
21496 51864 A 21624 45 277	
21497   51865   A   21625   171	
GHCCMNSSQNKI	1
HALNVKRFTIFQS	7
	-
QEVRCE*RDFPPE	EATWTGLPV
LPSQ	
21498 51866 A 21626 276 387	
21499 51867 A 21627 1 218	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21500	51868	A	21628	139	9009	ESSEDVSGGLPFLVDLGFRSVC
			į			VSLFHVKNPVQWDQKLPKKKK
						PVRSYCGSKLLQREVIQPDVME
						EMVVSCVIKHLNLVDALQSLIN
						FQYQEEHAEEYDLLCKIMGETF
						KKLNAMERQLQNKMKELELLC
						SMKEVSFDGNDLENMVLSLRE
						KFLQEVNSLIQKPSHPLAKTKT
						LVKSLMNRAELLLHVTIAAQSG
			1			LTRSISGTPAETPACKSASETKV
						ISHAVRQPVFLRSMSAPSDLEMI
		_				GNEDLEFTRANQRR
21501	51869	A	21629	1697	2056	W DENIK W OD TOD NO TWO TO CAR A
21502	51870	A	21630	237	1545	KPENKKSRTGRNRTKSTPGYRA
						/RPAAATARISARSTK*WKNSL
				1		AGRWARHICWTLWALIPRITLR
			[			/HVMAAGFPQRMQKDYRDAID ALCDANRFGQKKRLGFWRYKE
1						DSKGVGLCSYLLIGFYYTDPKN
						GAAAMKAFVVTRVGDVFLAFA
					1	LFILYNELGTLNFREMVELAPA
						HFADGNNMLMWATLMLLGGA
						VGKSAQLPLQTWLADAMAGPT
						PVSALIHAATMVTAGVYLIART
						HGLFLMTPEVLHLVGIVGAVTL
					1	LLAGFAALVQTDIKRVLAYST
					1	MSQIWLRCSSRLACRHGCGGSL
						ECLQHEGIPYSRMGRPHTTSAL
						RRFTSEFGMGSACFTLGYFFMY
						KQAVILLLMLFTASVSAACTLPI
					1	AIDHGTPGPAELLTVALRYNLS
					1	SYDAAYLELALRLEIPIAKGSPN
						FAFRRAATGNEGIKMHFLRNK
	ļ. <u></u>					GILWI
21503	51871	В	21631	1	1440	DVTD A CDVTD AVOI TO COVD
21504	51872	A	21632	525	716	PKTPAGPVEPMHQVPSSKRVPT
						RDKRTRGDSTRHSGQSKPTKRY
21505	51072	1_	21622	0	126	NHAYEI*PRIKWFIPDAGC
21505	51873	A	21633	1476	136 1611	GCDPAPGLPAPPHT/QP/DH*NQ
21506	51874	A	21634	1470	1011	ROIGIG*FLTGTVPATHLRORLA
						AA
21507	51875	A	21635	461	827	AALQMRDTSDPADCGEQSVCG
21307	31073		21033	101	027	YSEPVPEYACFHRIQDLHDTET
						AACFSHAGSGRTHV*SQSEHAA
						GSVLWSYRYRC*TGTAGYTDR
						FSQQRCHVRRGISPRCWPEYQS
						RSWCRLFPSPESAG
L	<u> </u>	<u> </u>	L	<u> </u>	<u> </u>	

			SEQ ID NO:	Nucleotide		Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
21508	51876	Α	21636	1	3735	MAEASVFQTVDPWRPGFATIPD
ŀ						FLKERYDKTTRIIIDFCFLIATGV
						CFLPIVLYSGALALNSLFHVGES
		ļ				LQISHGAAIWLLVILLGLAGILY
						AVIGGLRAMAVADSINGIGLVI
						GGLMVPVFGLIAMGKGSFMQG
		İ				IEQLTTVHAEKLNSIGGPTDPLP
						IGAAFTGLILVNTFYWCTNQGI
						VQRTLASKSLAEGQKGALLTA
						VLKMLDPLGLKPDLLAYNTAL
						RALIPQRLIAFDDAVRSTLKEEE
						KLVNSSDWGY
21509	51877	Α	21637	840	1064	WSVENYHRRGYGGHIPLPAASP
						AVAAQWCDAGR*VPPAARRW
	ļ					HRHQVYLTYLITLDRGLSRARF
	Ì					CIPSAHCSAY
21510	51878	Α	21638	979	1409	RLVVDIFHDNRHAFPRRLVSDA
:						AAHNACTQHRGLFRGFNVFGE
						FLRFAFYILIVKENADQCAGFV
						GMRQRDKTFVFQV*GFFTTKT
		İ				GRSFNGFYCGDGGRIMF/LPLFV
		ł				RPSL*RW*SSWWLQSCSVLVVP
						ASADVLLSSRGRR
21511	51879	A	21639	903	1397	LPTLPMMVCCTTPTSSRKHWM
						*NPSTLT/HAWVPYTNFSPIYEG
	1		;			KCGMSGARVEGKVIYETQSTH
						KLLAAFSQASMIHVKGDVNEE
						TFNEAYMMHTAASPHYGIVAS
						TETAAAMMKGNAGKRLINGSI
						ERAIKFRKEIKRLRTESDGWFF
						DNAAGYIDTITLGGPS
21512	51880	В	21640	1	5305	
21513	51881	Α	21641	129	723	LITARGQVADRNLRDLLAPWV
						PDAPSRALPEMTLDSRVAAAV
						DLFVARIGHQADGRRYIPQAIA
						QGVAAIIAEAKDEATDGEIREM
						HGVPDIYLSQLNELLSALAGRC
						YHEPSDNLRLVGVTGTNGKTT
			ł			TTQLLAQWSQLLGEISAVMGT
						VGNGLLGKKKEKAAIQYNTLC
			}			KSCTLTHPKPHNESDH/IRCRC*

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21514	51882	A	21642		2230	MMRTSGGTRDNESSSDDERER ENRLSDEKSNEYANLVGGERY EPDEENPMLGFRGAGRYVSDSF LRLCSSLEIVKARETVALIQGLK H\DLLADDTALRALIPQRADRFP MNAVRSTLKEEEKLVNSSDWG Y\DAQAFARWRPEYGYFAKQA GFTVKTSASLAALWQVVNQIG GKERYFFGNILWQTRALMDRAI GHKLAKGRPEREYLQTGDAVD SWKVIVVEPEKQLTLLFGMKAP GLGRLCFSLEDKGDYRTIDVRA FWHPHGNRAEREKPRFQPYSRR KSFIGQNMLLPMSIMPPVFRRVI CPLGVARLGKFRDPVYLVCVFT RQIFRYYTSGDEFLQINTTGGED IDDIDDIKLFVYEESYGISKESH WREAINAKAMGAMTLNWQEK RWQRFFNSEEPGNIEPVYMLEK VENQNHAKWEVHNFTMGYQR QVTEDTYEYLLLNVVGVPVPD NIHPMTKDEAGVWSWRTPILK GNLYEYFFNVDGVRSIDTGTA MTNPQRQVNSSMILVPGSYLDT RSVAHGDLIAITYHSNALQSER QMYVWTPPGYTGMGEPLPVLY FYHGFGDTGRSAIDQGRIPQIM DNLLAEGKIKPMLVVIPDTETD AKGIIPEDFVPQERRKVFYPLNA
			i			KAADRELMNDIIPLISKRFNVR KDADGRALAGLSQGGYQALVS GMNHLESFGWLATFSGVTTTT

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21515	51883	A	21643	<u> </u>	11549	MDASGHHVRGFASGMGCCRV
21313	31863	^	21043	1	1349	RRTMKRETGGGRLAGFRLVGT
		l			1	
						LGQHLGAHRSASKGHQILAAA
					1	RHVDRLAK\LQLA\NVSCHEVD
						LSWPDNLPALLQDIDTVYFLVH
						SMGEGGDFIAQE\RQVALNVRD
						ALREVPVKQLIFLSSLQAPPHEQ
						SDHLRARQATQ\EIRSEGNVP\V
						TELRAEIIVGAGS\AAL\EVM\RD
						MVYNLPVLTPPRWVRSRTTPIA
						LENFLHYLVALLDHPASEHRIF
						EA\AGPEVFSYQQQFEHFMAVS
						GKR\RWFDPPSPLPT\RWISVWF
						LNVITSVPPTTARALI\QGLKHD
						LLADDTALRALIPQRLIAFDDA
						VRSTLKEEEKLVNSSDWGYDA
						QAFARWRPEYGYFAKQAGFTV
						KTSASLAALWQVVNQIGGKER
l						YFFGNILWQTRALMDRAIGHK
		:				LAKGRPEREYLQTGDAVDSWK
ŀ					•	MIVVEPEQQRTL\LFGMQAPGL
Ì						GRL/CCFSLEDKGDY\RTID\VRA
						FWHPHGMPG/L*F*WLLMIPAH
						LFIFRGMAKQIARLAVPSTDLID
21516	51884	A	21644	201	363	SLRGSRTAASPCWAVPVHVPG
						НGАGЕРАННРGНQР*LРРРНРН
		1				VLLPLQPVLA
21517	51885	A	21645	3	440	PFSLPFSLGLTASLFMNTWLAIS
						SNCDGFIPASPACFCSFLLMSSA
ł		1				L*RTKAM*IMC*FSGLSGSKGV
		1	!			YIR*ASHSLS*NCAGLSSFP*MT
ł						SESLLTFVAF*APLINPSKRASLS
						RFSLCISSLSCGSNWGSVPGNW
İ		İ				VLPYSWGF
21518	51886	A	21646	359	543	VVTPDSGSSH*KANSFCLSGAN
21310	31000	``	210-10		343	LMLKKASFRSKQVNQLSSAGS
						NPSNMYGLGTVGCKVTVA
21519	51887	A	21647	2	582	THE SHAFT OLD I VOCK VIVA
21520	51888	A	21648	3	574	
21521	51889	A	21649	1	195	LCVFFVLVSFFLSPLDSQLAVV
						C*FDGTGIGMYLTSAVSPPPRN
						GVVASVMYAVVTPMLNLFICS
21522	51890	A	21650	2	457	
21523	51891	A	21651	21	545	PPCTSLRPLHAFSGKMTHLNRG
		l.,		<u> </u>		TLSLLDSQLHNLIALQMTCFKD
						VEIPNFFWEPSVTPSHRNINMYF
						PAAVFGFLPISGTLFSYCKIVSSI
						LRVSSSGGKYK/AFTTCGSHLS
		1				VVC*FYGTGVGGYLGSDVSSSP
	1					RKRAVASVMYTVVTPMLNPFI
L	l	<u> </u>	L	<u> </u>		YSLRNRDMKSVLRRPHSSAV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21524	51892	A	21652	899	1039	LWTSSLTAESSPMRAA*LRETSI
21324	31072		21032			VSAASRAFSAALSSSALRAFSA C
21525	51893	Α	21653	804	1390	QGTIILKVLRHLRWTWIPSRASL
	ŀ					RTGYCFCLTGWTPGLTSISTGA
						WLTANPGGLSSAHTLGHRLAR
						AGLISWPGSVKKSLHSSSRGSIR
						VIMTPVPGNFSSKEPCSVKEIVA
						LSLLSKSLPNKGKGQSGMYKN*
						*ITLCPPIVQVWGKQKACFAET
						PMAPMMSIVFLDKGATEAVTT
			ļ			ECSALVSTRKSMSLPPTV
21526	51894	Α	21654	84	730	HLSDSKSQVRYQPSADLSFLLG
						HYNPQSSRRVLSVLDSQLHNLI
						ALQVTCFKDVEIPNFFCDPSQLP
						HLACCDTFTNKIIMYFPAAIFGF
						LPISGTLFSYSRIVSSILRVSSSG
						GKYKAFSTCGSHLSVVC*VYGT
						GVGGYLSSDDVSSSPRKAAVAS
1	İ					VMYTVITSMLNPFIYSLRNRDIK
						GVLRQPHGSTVQFQYLLICSIPF
						VVWVKKGSKVK
21527	51895	A	21655	1049	1979	
21528	51896	Α	21656	2244	2901	LRCLSLPFLEAWKRDMLLSVM
	1					A\YDRFVAICHPLYHSAIMNLCS
:	-					VGFLVLLSFFFF\LSLSDSQLHN
						LIALQMTCFKDVEIPNFFWEPS
						QLSHLACCDTFTINIIMYFPTAIF
	1					GFLPISGTLFSYYKIVSSILRVSS
						SGGKYKAFSTCGSHLSVVC*FY
						GRGVGGYLSSDVSSSPRKGAV
						AAVMYTVVTSMLNPFIYRLGN
				<u> </u>		RDIKSVLRRPTEHCLISKK
21529	51897	Α	21657	1	2850	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	1	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21530	51898	ĪA.	21658	Ti Ti	1703	KKMALTSFLPAPTQLSQDQL\E
						AEEKARSQRSRQPSLVSSRREPP
						PYGY\RKAWIPRLLEDF*DRGA
						FPEIHVAQYP\LDMGRKKKMSN
1		1				ALAIQVDSEGKIKYDAIARQGQ
		1				SKDKVIYSKYTDLVPKEVMNA
		1				DDP\DWQRP\DEEAIIEITEKA\R
	1	1	1			VAL\EKSVSQKVA\AAMPVRAA
	1	1				*QIGPLLQYIPIHTISGKGVAIQT
						SWS*TRRVIR\MVEMQ\KDPM\E
1						PP\RFKINKKIPRGP\PSPPAPVM
						HSPSRKMTVKEQQEW\KIPPCIS
						*LGKMAKGY\TIPL\DKRLAAD
					:	GEEGLQDSYTLNGKFSAKLAE
						ALYIADRKAREAVEMRAQVER
						KMAQKEKEKHEEKLREMAQK
		İ				ARERRAGIKTHVEKVKYFSTVH
						TEDGEARERDEIRHDRRKERQH
				İ		DRNLSRAAPDKRSKLQRNENR
					:	DISEVIALGVPNPRTSNEVQYD
			}			QRLFNQSKGMDSGFAGGEDEI
						YNVYDQ\ALIGGKYMAQSIYRP
						SKN\LDKDMYGDD\LEARIKTN
						RFVPDKGVFLVQTRRQRGREGP
						\VQFEEDPF\GLDKF\LEESPTAM
,						GGS\KRPSG*PAAPREHGAWKA
						RKRRKE
21531	51899	A	21659	2	1588	CLRRALPGSIESAPTAPPPPPSGE
						GSQAYISRLPATLIVLILERRQL
						LGVWRRARCCSLINLAHRRHC
						GMPTRESPCAILARRYEGRRSPS
						TVAQPPPPPPPPIELAPSTSETPIQ
						ATRGAHPAQPSS/VPPA/PARPPE RPPVPEQGPSGTHPDPTP/PGPT
						NAPQRPSAP\PNTTN*PTRHMH
						QLPDAA*TPYPAYGSDYL*AR*
						DATASHQASCTN/SPDAA*TPYP
						AYAKIERA*ENPVPSAIKIIGRSS
						SSGRQASPWGMA/SPHWRSWR
						MGPNGSGKRPRRAAERISCCRN
						AQGRTHSQKAYIQQQPQCCVN
						VHLQPPD/PTPAQ/PSQTLSPPDS
		1				/PAPPLPAYPS/TQHPTAPPPTSSP
						MSS/RIVKPHPQHCGARPGQGK
						ERDETERKDAKERNRTATPTT
						RHTDRLPRTAHGPPPP\PPPPPP
						PPPPPPPRDSPRRPRREKPEQQ
						KEGGQTGRTQRKETARRGEER
						ETGRKKTRGAPPPRPAAARRPH
						ETAETKRRPPARQHRPPRRRPS
			1			DGTERTRHQLGFIMLGLLSWN
L	<u> </u>	1	L	<u> </u>	1	DO LEKT KINGEOF IMPOULES WIN

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		_ · · · ·
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21532	51900	A	21660	2	192	RVDICSTNYCWWLYPMTSGPSF
		i				SYV*GP*EEDVRE*MCGRDGL*
						YLSIGVVCGFICQHHNRAQ
21533	51901	Α	21661	783	2119	GSEMPVTVMLTLGEPGHFMSS
						LG\CWCGVEDEAAPSKQSIYIQ
						RETQVRTPMAGVSPKKAHPCE
						MCGPILGDILHVADHQGTHHK
						QKLHRCEAWVNILYDSGNFHR
						RQNEHIGEKPYRGSVEEALFAN
						RCKLHVSG\SHLSSVRV/EKDFL
						LRSGLLQQEATHTGKSNSKTEC
						VSLFHGGKSHYSCGGCMKHFS
-						TKDILSQHERLLPTEEHFVWCE
	ļ					CGKSSSKYDSFSNHQGVHTREK
						PYTCGICGKLFSSKSHLLVHQRI
E	İ					HTGEKPYECEVCQKFFRHKYH
						LIAHQRVHTGERPYECSDCGKS
						FTHSSTFRVHKRVHTGQKPYEC
						SECGKSFAESSSLTKHRRVHTG
						EKPYGCSECEKKFRQISSLRHH
	:					QRVH/T/NERAYECSESSLIKYW
		.]				RIHT*VRSCDCSKCGK/CFTQRS
1						ALLGYWRVHTR\KSPYK*NKFG
1						QFCSHTSVFLQDYS
21534	51902	A	21662	84	437	KMTSTQPVKFSLFASHSVPCGQ
						SFPGRRAFAPAMPRRRIPSSDHL
		1				PEPGQSHQHVNRQRWACRELI
						AAFLPSSYRL*LVETGGEIGQRN
		1				RASCHSPFPAGSSPGIFQTEIGKP
						CLLP
21535	51903	Α	21663	69	446	
21536	51904	Α	21664	880	1365	
21537	51905	Α	21665	1705	2196	RQNQREGDDDTEDQPPRPGTQ
						MTELFFLIQMLCFMANALRQL
						VHPL/MGNVHEYFADLKMPAC
						AVSAFLCAMHRSITSMFE*DNG
					:	PGPADADAR\LDMRIQVFMFCA
						NTLPARVLQRNKNPGQDHPET
		ŀ				AERRAESQRIIEKGLKCIQGKGI
						VNSISMKEGVDAFIH
21538	51906	Α	21666	1	1147	
21539	51907	Α	21667	254	557	AADEIGGIAVETVTEDAHLLCG
						CTVVAIPPRICVSAGGGAGDRK
		-	}			SSVRFHQNLAQVHITRGV/VWT
			1		1	MTNSHIITCCATAFGTTTSPSPT
						ESTGVPCGRRNQH

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	l e	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
21540	51908	Α	21668	557	1031	SSRNASGAVRKISRGMPDKKW
						NMALT*HSR*ANFSPLPVSGLS
						RRKICTIPRAQRIR*PICADRLSV
						ASPAACGI\AYTRRYSHDGATA
						EKYAHLQSRFRQQCRQFHPAA
						YDESPHRNRRRKWRPTYRCHP
						APDHRTACLRSAFYGNAPGCV
						QTERSW
21541	51909	Α	21669	572	825	QARISTSFFRESLPSRHPVPRYR
						PLEAFWTPCTRGYYRFFFSPAR
						ENAEYRTL\KHWAKFLLVRASF
						PGASTLCCKSPGRTISV
21542	51910	Α	21670	303	899	EDNYSGSSLYCFDDVASVQMP
						GSNGNCRR*CAGKTSG/TWRNS
	ļ					LVRQWLLTAQKKTLLHAVSNL
						PKTWAQILFSKQRVLRSPLNRH
						LIW*CAAVKL*LLVLYPAIRQSI
						SSKSIAKSLSRRYFR\LPIGDSM*
						NRW*RIFTIIGMYIGTAL*SDGAI
						VVCRLCENLPGSFHQRHQRCD
						AASLAGGSEPIAFSRAGRTPGP
21543	51911	A	21671	1	1320	
21544	51912	Α	21672	99	761	IDNAPPDQTTVHAFAANSENHS
						PPQLPYETPDDADPTHQAFSRW
						RHRPQAYSADALLRERLPPTHR
						HRGQSPDSLTSLPYWRGRRKY
·						RAAPALT*TPPDTLFDSPP*PAV
ļ						PPPVAKELPVTAQRQVHNAMS
						ANSFVIFHPVKIRNNIVLQPAIAI
:						QRRNGHGVTRCAAALHSRHGS
						LSVARHRVHLPVCRQADDKPA
						PAPLRPARYCLCPNQSAAVRYL
						R
21545	51913	Α	21673	62	680	SSRNASGAVRKISRGMPDKKW
						NMALT*HSR*ANFSPLPVSGLS
						RRKICTIPRAQRIR*PICADRLSV
						ASPAACGI\AYTRRYSHDGATA
						EKYAHLQSRFRQQCRQFHPAA
						YDESPHRNRRRKWRPTYRCHP
						APDHRTACLRSAFYGNAPGCV
						QTGPVRRSGAASASSPAVFLSG
						TSPWSFARTIALVRGRSRKSTRT
						RLWHISAHC
21546	51914	Α	21674	411	481	

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
21547	51915	Α	21675	402	915	RLASKLFTTVTPRGGVGELQAT
						TRNAAVAGIKRKRSNRGRPAEE
						PELTADRGQGENEITEAATAAK
						ARRTMRPKTGNQNPDKHRTRG
						LQICHTFNANGNRARCSAANA
	ļ					MAATISGLSSGLSGTGWQETIR
						PPLAAIF*SSCASNDIHLPKSFLA
		ļ				LPNGTAPLNGLPSIAS
21548	51916	Α	21676	1416	1832	GSAPRQRHLAGRRPVTTRPKHV
,						SHHGPQENSAHRLSHVDNESTC
						IVCGTLRLQHSARYFPPSLPETL
						FLAPVNHSV/EYNWQAAVPK*I
						Y*HQTFPGRDCQHLLFHFVRKF
						GVRESI/AQVVAVRAATSAWRY ISRSSAITL
21549	51917	A	21677	97	300	SKSSAIIL
21550	51917	A	21678	756	1179	RRRLPSPYCRAAAGSSPSYECF
21330	31710	1	21070	,,,,,		AHQRSVASSRGHTGTRQFYRE
				ı		HWFLRCIRRLSGTRLLAYHRCF
						QPHLACSGKRAVSASSTRSADV
						AAWYRV/CAKPFSSSS*VQEDL
						SYLRFSAS*P*TVIL*PACRKPCR
						KRRGQRRWR
21551	51919	Α	21679	3	1279	
21552	51920	Α	21680	1	2448	
21553	51921	Α	21681	317	1985	PNTPLTSKLPLLTHPLTLLTPSA
						SGWQKTTKLSCVF/PETEKYAH
	1					ERLAHHVAQGANQYAPMTGV
	}					QALREAIAQKTERLYGYQPDA
						DSDITVTAGATEALYAAITALV RNGDEVICFDPSYDSYAPAIALS
				l		GGIVKRMALQPPHFRVDWQEF
	1			İ		AALLSERTRLVILNTPHNPSATV
						WQQADFAALWQAIAGHEIFVIS
				İ		DEVYEHINFSQQGHASVLAHPQ
						LRERAVAVSSFGKT\YHMTGW
						KVGYCVAPAPISAEIRKVHQYL
						TFSVNTPAQLALADMLRAEPEH
						YLALPDFYRQKRDILVNALNES
						RLEILPCEGTYFLLVDYSALSTL
						DDVEFC*SLTQEHGVAAIPLSVF
						CADPFPHKLIRLIQTAAINLLVN
						HERRANAVINVDHAEIPFLRIIL
			1			PFLRQRQHHAVIIDHHPNAELL
						FQQIFHLHRRCPRRFRHGLRHA
						GHRINLTVDFQRQVDGMVRPQ
						FTVAMPAIFRFRTAAPSASVHQ
					1	SGVRLPYQFMALIASDNFCCW
						VHGGYITLAVNGNDAIGSVFNN
					1	QIVKCPRFTQCLGEVINTLTDLP
	1	1				HFTGQLW

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	1	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21554	51922	В	21682	160	312	
21555	51923	Α	21683	80	452	
21556	51924	A	21684	158	1980	REDTAHLPAVGFLPQNPFRAIR
21330	3172.	` `	2.00.	1.50		ASGEMLLPLLLLLPMCWAVEV
		İ				KRPRCV\SLTNHHFYR\ESKPFT\
		ļ				CLDGSATIPFDQGQRLTYCDCK
1		į				\DGS*RSQATAA\CP\NGQAFH\C
				İ		HQTLGL*SPL**SPSKPGSNDGV
						CDCCDGTDE\YN\SGVI\CENTC
						KEEGPVRRRESLQQMA\EVTRE
		İ				G\FRLEEDPLFEGLGKKAREEK
						QKKL\IELQAGK\KSLEDQ\VEM
						LRTVKEEAEKPEREAKEQHQK
						LWE\EQLAGFQGPNREQDARA
						ADAFKELDDDMDGTVSVTELQ
						THPELDTDGDGALSEAEAQALL
						SGDTQTDATSF\YDRVWAAIRD
	ļ					KYRSEALPT\DLPAPSAPDLTEP
		•				KEEQPPVPSSPTEEEEEEEEEE
						EEAEEEEEEDSEVQGEQPKEA
1						PPPL\SPPQPASPA\EEDKMPPYD
						EQTQA\FIDAAQEARNKFEEAE
						RSLKDMEESIRN\LEQEISFDFGP
						NGDV/FAYLYSQ/CARDLTTNE
1			ļ			YV/YTRLCPFQVLFFAGNPNSG
						VFPTSLGHLGASW/VLGPDHGQ
						VSVAMKF*ARHGACWQGPNPL
						QHP*RFWFGKET\MVTKHPQ*P
						SRCEYLHGS**TPAA\CPEPPAL
						KQPQPKD\DHDEVLSWMGARE
21557	51925	A	21685	59	359	LLRIGNDWCIPKGITPTTHIIKLP
21337	31723		21005			IGEIROPNATLDLSQSVDNEYY
		ĺ				CLLLAKELGLNVPDAEIIKAGN/
		l				GARVSGRTF*QALEC*ANGFTS
		-				LATGGYVSDI
21558	51926	A	21686	1	213	MSIAVTTTDPTNTGVSTTSSSSL
21336	31720	\\ \hat{\chi}	21000	['		TGSNAADLQSSFLTLLVAQLKN
ł						QDPTNPMENNELTSQLAQISTV
					,	SGIEKLNTTLGSISGQIDNSQSL
						QASNLIGHGVMIPGTTVLAGTG
		İ				SEEGAVTTTTPFGVELQQAADK
						VTATITOKNGAVVRTIDIGELT
						AGVHSFTWDGTLTDGSTAPNG
		1				SYNVAISASNGGTQLVAQPLQF
						ALVQGVIRGNSDITTFGVNMKL
		1				AHLGRQALMGVMAVALVAGM
						SVKSFADEGLLNKVKERGTLLV
						GLEGTYPPFSFQGDDGKLTGFE
		1			[	VEFAQQ*FAHGQQRRRFTKQFS
						DFAGGAAEKPGPDQSNGKQRA
		1				DVAIGTNQHGQRD
L	<u> </u>	<u> </u>	<u> </u>	<u> </u>	L	DA VIO LIIÁUOÁVO

NO: of peptide hod in USSN location of first		Amino acid sequence ( X=Unknown,
	1	*=Stop codon, /=possible nucleotide
sequence 09/540,217 codon for pepti	ide of peptide sequence	deletion, \=possible nucleotide insertion)
sequence		
21559 51927 A 21687 1	2339	MSGGQTEKQAQLIRDFQPDMI
		MVTPSYCLNLIEELERQLGGDA
		SGCSLRVGVFGAEPWTQAMRK
		EIERRLGITALDIYGLSEVMGPG
		VAMECLETTDGPTIWEDHFYPE
		IVNPHDGTPLADGEHGELLFTT
		LTKEALPVIRYRTRDLTRLLPGT
		ARTMRRMDRISGRSDDMLVLR
		RRHLCRPSACQLCRPPSLPATD
		RVLELFAQSGAQWLVILGDVL
		NHGPRNALPEGYAPAKVAERL
		NEVAHKVIAVRGNCDSEVDQM
		LLHFPITAPWQQVLLEKQRLFL
		THGHLFGPENLPALNQNDVLV
		YGHTHLPVAEQRGEIFHFNPGS
		VSIPKGGNPASYGMLDNDVLS
		VIALNDQSIIAQVAINPAPEKEG
		FLIMEQRRLASTEWVDIVNEEN
		EVIAQASREQMRAQCLRHRAT
		YIVVHDGMGKILVQRRTETKDF
		LPGICATSKVRKLLCKSAALLP
		VSELLLVLLTPVFVGSAVIGKL
		GAFAIHIDGITVKRRLVRNTVL
		QFCGRSRQPLEYVSAGDTGPVL
		RALKRMMAMRHYMRSQTVKG
		VTDTRAIDEVALSVAQVEEMY
		RYLAIANYEDRFVIPTSHREMA
		GDAFAERNGCGFTFGDGCHGS
		DSKFNLFNSSRIDAINITEILKVI
		GLLMEYPDELLWECKEDALALI
		RRDAPMLTDFTHNLLNAPLLD
		KQAEWCEVFDRGRTTSLLLFEH
21560   51928   A   21688   391	631	
21561 51929 A 21689 2779	3348	LMQLHNGLPVAGLRQFCQQLV
		IQRLYFFTTATAHQCTRHMCHR
		HRSLLTAHRQMALCFQYVDDL
		HHAFADVLINNACYPLLMGEQ
		GVFIYNGTQQLLHWVIIDSRQD
		HGFGGL*KSRIVCERRRL*SQH
		GQFVFRHHVPDHRIDIAFNNAL
		*TLVIREISSVPASVGVGIDFPGG
		AATWAGPVVIKPGQS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
		ĺ		sequence		
21562	51930	Α	21690	1	1239	MPNVDVVGEMVNTMSASRSY
						QANVEVLNTEKAMSIAVTTTDP
					:	TNTGVSTTSSSSLTGEQRRRFTK
						QFSDFAGGAAEKPRPDQSNGK
		ŀ				QRADVAIGTNQHDAARILRKG
						TMTTLLNPYFGEFGGMYVPQIL
		l				MPALRQLEEAFVSAQKDPEFQ
						AQFNDLLKNYAGRPTALTKCQ
						NITAGTNTTLYLKREDLLHGGA
		l				HKTNQVLGQALLAKRMGKTEII
						AETGAGQHGVASALASALLGL
						KCRIYMGAKDVERQSPNVFRM
						RLMGAEVIPVHSGSATLKDAC
						NEALRDWSGSYETAHYMLGTA
						AGPHPYPTIVREFQRMIGEETK
						AQILEREGRLPDAVIACVGGGS
						NAIGMFADFINETN/VSVGPQH
						AYLNSSVRAD*VSITDDEDLQA
						FKTLTASYKTTPAAAKTSLHVP
21563	51931	Α	21691	32	1778	WDARVFSHSGWVLYLIRQYCSI
						SAVSSCGSTTGSACSATSWATA
		l				TSDSVTHLRLFSYRGSDHFRMR
						RFNHFRLRFGNLNGGFYDTGHF
						FNNWRANNRFDSGGDRDLSHH
		İ				RLHMYFFLFALLFNLYILRTYN
						WIADPDFARGQFRRRARYGQR
						HWRLGWITLVTIATTTLTTYAQ
						VTRRAARTTRHATVVTVISIVL
		ĺ				FLLFWCSLRYDWHYCILILRSTP
				ļ		SVLSQYPVGNELRRRRLVHLHL
						LCTYSDDDSRDPRYRQESRTRN GVVPVIVNIDVHRFGQTGDQH
		1				GAMIIPVIFLLNVAMHERASGHI
						PNGNRTYYLSRSQPPVFALMVE
						LFEEDGVAVARYLDHLKMEYA
						FWMDGAESLIPNQAYRHVVRM
						PDGSLLH\RYWCDCHTSCD*SW
						LEDVETAKHSGRPPNEVYRDLR
						AGAASGWDYSSRWLRDTGRLA
						SIRTTQFIPIDLNAFLFKLESAIA
						NISALKGEKETEALFRQKASAR
						PMREYETGEQWDKPNGWAPL
						QWMAIQGFKMYGDDLLGDEIA
						RSWLKTVNQFYLEQHKLIEKY
						HIADGVPREGGGGEYPL\QDGF
						GWTNGVVRRLIGLYGEP
L	L	L	l	L	L	OWING V VAILULIUEF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21564	51932	A	21692	1	2514	MKTLIARHKAGEHIGICSVCSA
						HPLVIEAALAFDRNSTRKVLIEA
						TSNQVNQFGGYTGMTPADFRE
						FVFTIADKVGFARERIILGGDHL
		1				GPNCWQQENADAAMEKSVEL
						AVVIDLWSRAVIGWSMSPRMT
						AQLACDALQMALWRRKRPRN
		1				VIVHTDRGGQYCSADYQAQLK
		İ				RHNLRGSMSAKGCCYDNACVE
						SFFHSLKVECIHGEHFISREIMR
						ATVFNYIECDYNRWRRHSWINS
						LLSQKRNTQGRIEDGRQH
21565	51933	A	21693	1	1527	EESQKKII QUIEEDUKQII
21566	51934	A	21694	792	1602	HCLSARRYELVCQPK*AFVR*L
						LR*IAVLR\HLNIQHLRRSHSLR
						HGRRHTRKGERGTINIVNGTPI
}						HERSRNIDNRRSPSPYIRKRGAH
1					İ	TELRSDHPLLINQHWHHFAGM
1						VAVTNGRIVAVIRHHDQAIIWL
						HRLNKPGKVTVNDLQTFGITCD
					į	VFCMTSEVGFLNVGANKRRRM
						GVKFVEGDIHNPLAIHQILPAH
						AIRIVDDICHLTHQHGWHACFI
						QFLWHLHFFIDAIAHMHDLETV
						VISARTRRIDTNKAPFFAFCGIE
21567	51935	A	21695	1	1530	LSQQTVNTLQAGHQTIHFFMG
					]	VIHGQRRPTHRRDVVIGHQWH
1			ŀ			CAVMSGTHSNAVIIEKTANIIGV
İ					1	VFTEIKSNNAKPVLWAIESNAA
1						ATDGEIVFDGEILKSACEINDSD
1						KKIEVALGHYNAEQFRNIGERS
						PKIPFTIPLVNCPMTGEIANILGS
						KPGTIFTMLRDTGGIKPHERKR
						AVAHLTLSEREEIRAGLSAKMS
1						IRAIGYWR*IASPSTFSRE\VQRK
						GGHR\YYKAVDANIRANRMAK
						RPKPCLLDQNVPLRKLVLEKLE
1		ŀ				MKWSPEQISGWLRRTKPRQKT
		1		ľ		LRISPETIYKTLYFRSREALHHL
						NIQHLRRSHSLRHGRRHTRKGE
		l	Ì			RGTINIVNGTPIHERSRNIDNRR
		1				SLGHWEGDLVSGTKNSHIATLV
		1				DRKSRYTIIVRLRGKDSVSVNQ
						ALTDKFLSLPSELRKSLTWDRG
						MELARHLEFTVSTGVKVYFCD
		ŀ				PQSPWQRGTNENTNGLIRQYFP
						KKTCLAQYTQHELDLVAAQLN
						NRPRKTLKFKTPKEIIERGVALT
21568	51936	A	21696	2	140	IITNSISRPGDPPMSDPHS*SPAS
		-	,	<u> </u>		SSAKTETKSLFNAFLV
L	<u> </u>	Ь	L		L	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	t .	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21569	51937	A	21697	1817	4083	KGLLTSFHYIKTPEVPNLCVEG
21309	31737	A	21097	1017	4003	GTRLQIWPACSSTGSGGTCWK
						APPQPPRPPTSSSPS/PAPPSEAF
						VLSAGAMQLEAGAYSLVIIKES
						KNNVGVDVIKRKHFCTAGGNV
						NQYNHYGKHCVLSCSGPASGA
:						QPLYRMAQPPSPPNMAAGIPPA
						ARISSSITSPLLKPSWLHLLAHP
		i				VLQAHLTEEGLGRSHHVQRGQ
						CLVSKSVFTNACWEMQNVKRF
						TDFIPLHKNTRGPEPLCRRRNO
			:			APNMACMQQHWQRRN
21570	51938	A	21698	247	281	KPGKQHSMLK*SALLPSAHGPT
21370	31736	<u> </u>	21070	247	201	S/CKEARCHPVQILPPELLF*MP
1						KKICPPLCYLWTVKAWETTFH
						AEV
21571	51939	A	21699	531	716	PRRLRAAPPDLGGFRLLPR/LPE
213/1		ļ'`	21057	, , ,	,10	ASGCSP*SQRLFQKAPSPAPCS
						WRRSRFCVKTCVLPMGGQ
21572	51940	Λ	21700	202	606	ITCLHAHVTWCQILPYLMLPAP
			:			LLRPLPTPLTVAGVGQSLPEIQL
						SVTLPLQPGRNQGSLESRHLVS
						WAEPP*DPAVSALPLQPGRNQG
						SLESRHLVSWAEPP*DPAVSAL
						PLQPGRNQGSLESRHLVSWAEP
21573	51941	Α	21701	444	1167	FRLHGHLERGPFYFLSPLAVQS
		ŀ				SQSCLRRFTTRYHLP\YVLNRK
ĺ						NSWHVAHIFRRHAKPEEQAPIY
						SHIHFTSDLDEVLNDPDVKLVV
						VCTHADSHFEYAKRALEAGKN
						VLVEKPFTPTLAQAKELFALAK
						SKGLTVTPYQNRRFDSCFLTAK
						KAIESGKLGEIVEVESHFDYYRP
						VAETKPGLPQDGAFYGLGALFT
						NQQGFFKSSLLSIFADSSSSVAG
						RSPHNLVKKRGGIVAISAFAG
21574	51942	A	21702	ī	668	

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21575	51943	Ā	21703	1	1308	MRVYMDGRLIVSTRQRKVLAA
						GPMLYAATRAPAFTMSGDNLV
		İ				RSHASIYMTSCHTLLGEQAGYA
i						VFGALLPAITLLHLNSACCIVYV
						SCLLADAGHRDSLQQAGDKWT
						TTDSYSSSCTSETLGSACVRYA
						HSSRNHVRFALSDLDEVLNDPD
						VKLVVVCTHADSHFEYAKRAL
						EAGKNVLVEKPFTPTLAQAKEL
1						FALAKSKGLTVTPYQNRRFDSC
						FLTAKKAIESGKLGEIVEVESHF
						DYYRPVAETKPGLPQDGAFYG
	1					LGVHTMDQIISLFGRPDHVRL*
						HPGALRNKSPIWTNPFEAQL\FY
						G\DLKAIVKTSHLVKIDYPKFIV
						HGKKGSFIKYGIDQQETSLKAN
						IMPGEPGFAADDSVGVLEYVN
						DEGVTVREEMKPEMGDYGRV
						YDALYQTITHGAPNYVKESEVL
						TNLEILERGFEQASPSTVTLAK
21576	51944	Α	21704	199	706	SHDVCRINADRQRDRNNRRQR
	ļ					RSLRVYKLGNQQQNNGIGPRIL
						FHHFTQYALNFRHMRTEHGFC
						HPCDTVHRHDGDHTGTEDFVF/
						WR*TRDWFWSG*\YQRPGQQH
						KHLNNQYHRQRFAAEGWTQV
						RISTKKGDYSDCPEENPHCQPV
						AGRYHYSLVGRVSNATLARLIR
21577	51945	A	21705	354	569	SRAVCWQVGTRLWAIFGVS*E
		ŀ				LQRHVLVVFAGFWIVKNSRNL
				•		FLVCRAEHKRSVVKGLLRQQG
		_				AGLLVPLS
21578	51946	Α	21706	373	462	
21579	51947	Α	21707	1	1293	

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
21580	51948	A	21708	1	1371	MCPIEETASSFGGKPLSMVLVIQ
						MFMLLTGALIIILTKTNPASISK
	1					NEVFRSGMIAIVAVYGIAWMA
]						ETMFAPACYGYYILPTYPSDLA
						AIQFDRSGTTHIGRFVINHSFILP
						GLIGVSVSCVFGWIFAAIKRDA
						AAGRAKENVIFHHFPFQSVKAD
1						IATNPFKRPGGAFIRITQTFRTV
						QTFRLSAYRLDFAGDRLRISTPR
				:		AKMRTAFKKDHLRQRRRCIRQ
				ļ		RAPPARHNLVGAVALPATVAG
				ļ		VNFTFSNVPLDSSVLSSLLTDFS
1						TAVGSIVMLAVIMGLMLAFDM
						GGPVNKVAYAFMLICVAQGVY
						TVVAIAAVGICIPPLGMGLATLI
						GRKNFSAEERETGKAALVMGC
						VGVTEGAIPFAAADPLRVIPSIM
						VGSVCGAVTAALVGAQCYAG
						WGGLIVLPVVEGKLGYIAAVA
						VGAVVT\LFV*TC*KVWRVKM
						GLRLMKKKTTWIWILKLI
21581	51949	A	21709	961	1116	
21582	51950	Α	21710	228	388	RRGRNRYPALQ*TMFLPADGP
						ARCLSRQAAILKESVLPK*GRP
						DVVFSPAYPY
21583	51951	A	21711	1168	1851	LHRTGQRPAAQRTETHHPFFNH
						FARLQVDTVVINHHQHTVAFQ
						NRTFRSEI*RNNVDIFQPDVLPD
j						ILLGPVREREDADAFAFVNLPV
						VVVPQLRALIFRIPAMEAVTER
						VNSLFSAGFLFVTARTTEGGIKT
						VFVQRLFQAFSFHNIGMFRAAV
						HEWINPHRHAFRVFMHQQFAA
						VGFSSTIAELIHLAEFPAGIDMQ
						QRERQRTWIKRLARQMQHDAG
						IFTDRVEH
21584	51952	Α	21712	1	2653	
21585	51953	В	21713	632	709	

SEQ ID			SEQ ID NO:		I .	Amino acid sequence ( X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
21586	51954	A	21714	1	1214	MHDQQVIDAYHPDLRRTRAAS
						QSIIPVDTKLAAGITRFFPQFND
	}					RFEAIAVRVPTINVTAIDLSVTV
	-					KKPVKTDVGGAGTVKLNRCQI
				ŀ		ARISRQDVITPPGCVFEKALTLL
				<u> </u>		IASGRNDTLARQNNHTKESLVE
						KPFRKILVIKMRYHGDMLLTTP
	İ	į				VISTLKQNYPDAKIDMLLYQDT
						IPILSENPEINALYGISNKGAGTF
						DKIKNVLSLIKTLRANNYDLVI
		1				NLTDQWMVALLVRCLPARMKI
						SQLYGHRQHGIWKKSFTHLAPI
	ŀ					HGTHIVERNLSVLEPLGITDFYT
						DTTMSYAEDCWKKMRRELDA
						LGVKDHYVVIQPTARQIFKCRD
						NDIISKVICMIRGSPTEGRWSVF
						FFYPADFTFVCPTELGD/GC*PL
						RRTAETGRRRIRSIYRYSLHPQS
21587	51955	Α	21715	25	1086	
21588	51956	Α	21716	1	1113	
21589	51957	Α	21717	80	430	RWLATHYLTVKLSLLPNLAVT
						CQPKREQNITSNESTDC*VIVTS
						TKSDSLYTVGMLALSVRAIRCP
						LYLLTGLIFVSKNDLWYCELQS
						HYTVVLLLFMRKRSRFQSNVQ
						RKLMTNF
21590	51958	Α	21718	1	2076	
21591	51959	Α	21719	2742	3109	VWTEGTGNASRARR/WKGNQL
		1				LPVSLVKRKTTLAPNTQTASPR
						ALADSLMQLARQVSRLEKRAV
						SATQLMAKAVKGQSSLLPVSL
						VKKKNHPGALIRKPPLPARWPI
		1	ľ			LLIQLARQVSRLETGQ

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21592	51960	Α	21720	174	1608	TAAKRKIPSASIRSIND*KY*VC
21372	31700		21,20		1000	KRT*CVGQLAA\ENADGTVAEY
						NGYHVVFALAGSPKDADDTSI
			:			YMFYQKVGDNSIDSWKNAGR
ŀ						VLMNPLMILVKIIKLSLTLACH
						ALLDCKTLTLTELGRNLPTKAR
						TKHNIKRIDRLLGNRHLHKERL
						AVYRWHASFICSGNTMPIVLVD
		į			ļ	WSDIREQKRLMVLRASVALHG
						RSVTLYEKAFPLSEQCSKKAHD
						QFLADLASILPSNTTPLIVSDAG
						FKVPWYKSVEKLGWYWLSRV
						RGKVQYADLGAENWKPISNLH
						DMSSSHSKTLGYKRLTKSNPIS
1						CQILLYKSRSKGRKNQRSTRTH
		ŀ				CHHPSPKIYSASAKEPWVLATN
						LPVEIRTPKQLVNIYSKRMQIEE
	:					TFRDLKSPAYGLGLRHSRTSSS
						ERFDIMLLIALMLQLTCWLAGV
						HAQKQGWDKHFQANTVRNRN
						VLSTVRLGMEVLRHSGYTITRE
ł						DLLVAATLLAQNLFTHGYALG
21593	51961	A	21721	1	3285	DELVAATEEAQNEI THOTAEO
21594	51962	A	21722	1095	1188	
21595	51963	A	21723	568	1263	
21596	51964	A	21724	2	1688	
21597	51965	A	21725	1456	2046	SKGKVAIVSSHQEVRASPRRSP
21337	31,705	1		1.53		LGSEERPGPHTSRQAAGQPHPG
İ						TCSSVHCQGSWGLQTQNSEVIH
						DGDVEAISVGEDAAPGPRVHG
					•	ADLGQEEEAFR*VVMGGLPDR
		ł				DSALAPVPEPMHRGWAVRAAE
						GAVQNDPVGGSERSWGAGSHS
						QPFSHRPAPHTRGLAQAQTEAA
						GEGAVPEVHRGAVLAGVRAPG
						WAQP
21598	51966	A	21726	335	595	AGLFKSRKVSSICIRFE*ILTSCL
[ ]		ľ`				GVRISTGKLVARIHGSFADAE*I
	[			1		L\WTGGDNESVS/MR*FFRPLER
						DLYNRIWHEIGLLLVSLL
21599	51967	A	21727	80	428	RWLATHYLTVKLSLLPNLAVT
213/3	1707	l'`	[ ]		.20	COPKREONITSNESTDC*VIVTS
1				1		TKSDSLYTVGMLALSVRAIRCP
						LYLLTGLIFVSKNGLWYCELQS
		1				HYTVVLLLFMRKRSRFQSNVQ
						RKLMTNF
21600	51968	A	21728	622	1281	INITERIAL INI
21000	121300	ΓΛ.	21/20	1022	1201	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21601	51969	A	21729	1066	1827	VPQAKGNNVVITSYMTNRGFF
						EDKKATFAPSFLMNIKGNKTSV
						VKNSILEQGQLTWLQVAKRAG
ļ		l				LGGGQSGRTVLRERVRIEIAST
						HIALQHAIVTGDAVGMDDIPQE
						ARQYRHNQAYAYSIQGDGAED
						DDERIAFASHTIQECRSQGPGG
		l				YPFKIYSSRKDRIPISQSLLESHF
						TVHEETAPDLETTG*PGCKDHG
		ľ	}			NLCC/LSSVKLRYSMYRIKAGL
					:	WMTHPG*RTYWV
21602	51970	Α	21730	1	1326	
21603	51971	A	21731	1	1971	
21604	51972	Α	21732	1	1626	
21605	51973	Α	21733	1	1810	
21606	51974	Α	21734	305	442	AGLFKSRKVSSICIRFE*ILTSCL
						GVRISTGKLVARIHGSFADAE
21607	51975	В	21735	1	1938	
21608	51976	Α	21736	531	669	AGLFKSRKVSSICIRFE*ILTSCL
						GVRISTGKLVARTHGSFADAE
21609	51977	Α	21737	80	428	RWLATPYLTVKLSLLPNLAVTC
1						QPKREQNITSNESTDC*VIVTST
						KSDSLYTVGMLALSVRAIRCPL
						YLLTGLISVSKNGLWYCELQSH
						YTVVLLLFMRKRSRFQSNVQR
						KLMTNF
21610	51978	Α	21738	10	426	
21611	51979	Α	21739	314	451	AGLFKSRKVSSICIRFE*ILTSCL
						GVRISTGKLVARTHGSFADAE
21612	51980	Α	21740	1	1139	MWGIGSRTPIYNKILVYSSPEVS
						CEKPKYVKSQTSTHTGFESHKR
						YIFYLRLVGKNLHVSAMGLGK
		1				SQERRKKALKSQPQQPADSFPS
				1		LPTVEGLTTSGEVAAGFWLSGE
		1				ALSSC*LLSASVHLLLACLGMG
						KCSCSGFWLYF
21613	51981	Α	21741	306	444	AGLFKSRKVSSICIRFE*ILTSCL
						GVRISTGKLVARTHGSFADAE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
21614	51982	Α	21742	189	1089	HHTAHCQ*CWL*S/DHGINPLRS
						WVGTG*V/GVRGKVQYADLGA
						ENWKPISNLHDMSSSHSKTLGY
}		ľ			·	KRLTKSNPISCQILLYKSRSKGR
						KNQRSTRTHCHHPSPKIYSASA
						KEPWVLATNLPVEIRTPKQLVN
		1				IYSKRMQIEETFRDLKSPAYGL
						GLRHSRTSSSERFDIMLLIALML
						QLTCWLAGVHAQKQGWDKHF
						QANTVRNRNGSLLTNWPSVPP
						TIKEEENSEEELAAATTSKEQEP
		1				IGTDLDAVRTPEPLEEFPKREDQ
						EGSPPETSLPYKWVVEAANLLI
						PAVGSSLSEALDLIES
21615	51983	Α	21743	1	1135	
21616	51984	Α	21744	80	430	RWLATPYLTVKLSLLPNLAVTC
			}			QPKREQNITSNESTDC*VIVTST
						KSDSLYTVGMLALSVRAIRCPL
						YLLTGLISVSKNGLWYCELQSH
						YTVVLLLFMRKRSRFQSNVQR
		1				KLMTNF
21617	51985	Α	21745	80	430	RWLATHYLTVKLSLLPNLAVT
						CQPKREQNITSNESTDC*VIVTS
		1				TKSDSLYTVGMLALSVRAIRCP
		1				LYLLTGLIFVSKNDLWYCELQS
						HYTVVLLLFMRKRSRFQSNVQ
		<u> </u>				RKLMTNF
21618	51986	Α	21746	773	1015	AGLFKSRKVSSICIRFE*ILTSCL
						GVRISTGKLVARTHGSFADASF
		1				STDLYHGTLKPASLTMSGVVLL
		$oldsymbol{ol}}}}}}}}}}}}}}}}}$				GRMLARSARNWS
21619	51987	Α	21747	391	527	AGLFKSRKVSSICIRFE*ILTSCL
		↓				GVRISTGKLVARTHGSFADAE
21620	51988	Α	21748	671	1030	AGLFKSRKVSSICIRFE*ILTSCL
						GVRISTGKLVARTHGSFADAE*I
						LGDGW*Q*VRVER*FFRPLERD
						LYNRILACPLINT*PTPVVRLKP
		1				*TVWSMGPVVGTPARPIRTSGA
		ļ				CWWCW
21621	51989	Α	21749	221	478	AEQDCTILPVLRHFTFFKSRKVS
		1				SICIRFE*ILTSCLGVRISTGKLV
						ARTHGSFADDQTLFFMTLAVS
01600	41000	<u> </u>	01550	500	0.40	GELWTPSPWRPNSIRTS
21622	51990	A	21750	590	940	RWLATHYLTVKLSLLPNLAVT
						CQPKREQNITSNESTDC*VIVTS
						TKSDSLYTVGMLALSVRAIRCP
						LYLLTGLIFVSKNDLWYCELQS
						HYTVVLLLFMRKRSRFQSNVQ
21622	71005	1	01751	<u> </u>	(12	RKLMTNF
21623	51991	Α	21751	473	613	AGLFKSRKVSSICIRFE*ILTSCL
L	<u> L.                                    </u>	<u> </u>	L	<u> </u>		GVRISTGKLVARIHGSFADADT

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21624	51992	A	21752	342	1459	SFLWRFRLRGGSERFCLAGIPSA SGV*NIPSASIRSIND*KY*VCKR T*CVGQLAA\ENADGTVAEYN GYHVVFALAGSPKDADDTSIY MFYQKVGDNSIDSWKNAGRVF KDSDKFDANDPILKDQTQEWS
						GSATFTSDGKIRLFYTDYSGKH YGKQSLTTAQVNVSKSDDTLKI NGVEDHKTIFDGDGKTYQNVQ QFIDEGNYTSGDNHTLRDPHYV EDKGHKYLVFEANTGTENGYQ
						GEESLFNKAYYGGGTNFFRKES QKLQQSAKKRDAELANGALGII ELNNDYTLKKVMKPLITSNTVT DEIERANVFKMNGKWYLFTDS RGSKMTIDAKSTLLESTEGSMT
21625	51002	_	21752	2004	2201	PPMTATEPFLRFLKGEIAQPKPH
21625	51993	A	21753 21754 21755	81	2201 1080 430	HHTAHCQ*CWL*S/DHGINPLRS WVGTG*V/GVRGKVQYADLGA ENWKPISNLHDMSSSHSKTLGY KRLTKSNPISCQILLYKSRSKGR KNQRSTRTHCHHPSPKIYSASA KEPWVLATNLPVEIRTPKQLVN IYSKRMQIEETFRDLKSPAYGL GLRHSRTSSSERFDIMLLIALML QLTCWLAGVHAQKQEHPLWSL ALVRRSPAQSPHVGTGSSGST GCSSQAPCMKSNNALIVILGTV TLDAVGIGLVMPVLPGLLRDIV HSDSIASHYGVLLALYALMQFL CAPVLGALSDRFGRRPVLLASL LGATIDYAIMATTPVLWIYPLV RWLATHYLTVKLSLLPNLAVT CQPKREQNITSNESTDC*VIVTS TKSDSLYTVGMLALSVRAIRCP LYLLTGLIFVSKNDLWYCELQS HYTVVLLLFMRKRSRFQSNVQ
21628	51996	Α	21756	1	593	RKLMTNF MGVNDLWQILEPVKQHIPLRNL GGKTIAVDLSLWVCEAHPVKK MMGSVMKKHLRDFADFGTTIK QDFRLLGQTSVDRLLQLSQGQ AAYPIPDDQVQSVLISMLLIAL MLHLTCGLAGVHAQKQGWDK HFQANTVRNRNVLSTVRLGME VLRHSGYTITREDSLVAATLLT QNLFTHGYVLGKL*GDLSVLSL
21629	51997	A	21757	410	547	AGLFKSRKVSSICIRFE*ILTSCL GVRISTGKLVARIHGSFADAE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
21630	51998	A	21758	719	814	AGLFKSRKVSSICIRFE*ILTSCL
					GVRIFNR	
21631	51999	A	21759	1	1596	
21632	52000	A	21760	80	430	RWLATHYLTVKLSLLPNLAVT
						CQPKREQNITSNESTDC*VIVTS
						TKSDSLYTVGMLALSVRAIRCP
						LYLLTGLIFVSKNDLWYCELQS
						HYTVVLLLFMRKRSRFQSNVQ
						RKLMTNF
21633	52001	Α	21761	1343	1480	AGLFKSRKVSSICIRFE*ILTSCL
						GVRISTGKLVARTHGSFADAE
21634	52002	Α	21762	2	151	- Landana di Landana d
21635	52003	Α	21763	258	396	AGLFKSRKVSSICIRFE*ILTSCL
						GVRISTGKLVARTHGSFADAE
21636	52004	Α	21764	1	3081	
21637	52005	Α	21765	449	586	AGLFKSRKVSSICIRFE*ILTSCL
						GVRISTGKLVARTHGSFADAE
21638	52006	A	21766	563	2042	
21639	52007	A	21767	1001	1093	
21640	52008	Α	21768	1814	1870	
21641	52009	Α	21769	2760	3723	HHTAHCQ*CWL*S/DHGINPLRS
					:	WVGTG*V/GVRGKVQYADLGA
						ENWKPISNLHDMSSSHSKTLGY
						KRLTKSNPISCQILLYKSRSKGR
						KNQRSTRTHCHHPSPKIYSASA
						KEPWVLATNLPVEIRTPKQLVN
						IYSKRMQIEETFRDLKSPAYGL
						GLRHSRTSSSERFDIMLLIALML
						QLTCWLAGVHAQKQGWDKHF
						QANTVRNRNVLSTVRLGMEVL
						RHSGYTITREDLLVAATLLAQN
						LFTHGLFGPSVDAPPPKRGFKR
						PKLPELRLQRFFGGVITPPLESY
						SLPSDPVHLFPPRRPPRQLTASP
						LRARAPYECSSCKDA
21642	52010	A	21770	860	3503	
21643	52011	Α	21771	4597	6030	, , , , , , , , , , , , , , , , , , ,
21644	52012	Α	21772	1513	3464	AGLFKSRKVSSICIRFE*ILTSCL
						GVRISTGKLVARTHGSFADAE
21645	52013	Α	21773	1	453	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21646	52014	A	21774		1526	MSSSHSKTLGYKRLTKSNPISC QILLYKSRSKGRKNQRSTRTHC HHPSPKIYSASAKEPWVLATNL PVEIRTPKQLVNIYSKRMQIEET FRDLKSPAYGLGLRHSRTSSSE RFDIMLLIALMLQLTCWLAGV HAQKQGWDKHFQANTVRNRN CKMATLKEKLIAPVAEE\EANS\ PNNKIT\VVGV\GQVGMA\CAISI LGKVSWPDEL\ALVGCFWEDK LKGGNDGSFQHGELIFFQTPKN CGQDKGLFWGPPNS*RFVSW*L AGSPVQPRRGRIRLKSWVPEK WLMVFQIPLFLKIV\KYSP*FAS* LVVSHPV\DIPTYVT\WETKVGL PKHRVIGSGCNLDSARFRYLMA EK\LGIHPS\SCHGW/IFWGEHG DSSVAWCWNGVNVAGVSLQE LNPEMGTDNDSENWKEVHKM VVESAYEVIKLKGYTNWAIGLS VADLIESMLKNLSRIHPVSTMV KGMYGIENEVFLSLPCILNARG LTSVINQKLKDDEVAQLKKSA
21647	52015	Α	21775	2	355	DTLWDIQKDLKDL
21648	52016	Α	21776	2	178	
21649	52017	A	21777	1	888	
21650	52018	С	21778	127	255	
21651	52019	Α	21779	1	1824	
21652	52020	A	21780	330	476	
21653	52021	Α	21781	452	500	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21654	52022	A	21782	3	1584	DAWAAVSRGDEPGWGSGSWR
21054	32022	``	21702		1504	LLQHWGVSCWSDPKRFSSALQ
		l				GYIITDWCAPFNKRIWKREQPE
		1				ETFPNTETNGEFGK\RPAEDME
						EEQAFKRSRNT\DEMVEL\RILL\
		1				QSKNA\GAVIGKGGKNIKALRT
						DYNASVSVP\DSSVPERILCISA
1						DIETIGEILKKIIPTLEEGLQLPSP
		l				TATSQLPLESDAVECLNYQHYK
						GSDFDCELRLLIHQSLGREGIIG
		1				GQRCLKSKELRGGTLQTTIKLF
		1				QECCPHSTDRVVLIGGKPDRFV\
		1				ECIKIILDLISESPIKGR\AQPYDP
		1				NFYGWKPMDYG\GFTMMFDD
						RRGRPVGFPMRGRGGFDRMPP
						GRGGRPMQPYRRDYDDMSPRR
						GPPPPPGRGGRGGSRARNLPL
					·	PPPPPPRGGDLMAYDRRGRPGD
1						RYDGMVGFSADETWDSAIDTW
						SPSEWQMAYEPQGGSGYDYSY
						AGGRGSYGDLGGPIITTQVTIPK
		1				DLAGSIIGKGGQRIKQIRHESGA
						SIKIDEPLEGSEDRIITITGTQDQI
						QNAQYLLQNSVKQYSGKFF
21655	52023	Α	21783	211	1635	KNMETEQPEETFPNTETNGEFG
		1				KRPAEDMEEEQAFIRSRNTDEM
						VELRILLQSKNA\GAVI\GKGGK
						NIKALRTDYNASVSVPDSSGPE
					!	RI\LSISADIETIGEILKKIIPTLEE
						GLQLPSPTATSQLPLESDAVECL
		ŀ				NYQHYKGSDFDCELRLLIHQSL
		ŀ				AGGIIGVKGAKIKELRENTQTTI
		ŀ				KLFQECCPHSTDRVVLIGGKPD RVV\ECIKIILDLISESPIKGRAQP
	:					YDPNFYDETYDYGGFTMMFDD
		ŀ				RRGRPVGFPMRGRGGFDRMPP
						GRGGRPMPPS\RRDYDDMSPRR
						GPPPPPPGRGG/RGLGSRARNLP
						LPPPPPRGGDLMAYYKRGRP\
		Ì				GDPYDGMVGFSADETWDSPID
						TWSP\SEWQMAYEPQGG\SG\Y
						DYSY/AQGGRGSYGDLGGPIITT
		1				QVTIPKDL/ALGSIYWKKGGQR\
			}			IKQIPS*VGELSIKIDEPL\EGSED
						\RIITITG\TQDQIQ\NAQYLLQ\N
						SVKQYADVEGF
21656	52024	A	21784	860	1201	TISEPPALILLPIAVPSFSTTCRLP
		1				EDIIVVSAFPPLLIFCVPLLILVP
						VAMP*LLIFCVPPLIMVFVAFPA
						TSITPPLSIRVASALALVKTDIVP
						LSLIIVLFAEPYAFMSK
L	<u> </u>		J	·	<u> </u>	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21657	52025	В	21785	1	3750	
21658	52026	A	21786	1	1350	
21659	52027	Α	21787	189	1939	TAMAGRMPCAEVISGSATRPD
						GPKALFKPIGRPHQAIRPINKPP
		1				AAPPKQAVTNGFMRGRVPPKT
						LTVPSESSLAPMTLVVIVVKEA
						LPPAKIFSTPVPYQPTRRAPDAE
						PLSASKTPSVYLVNFVVSPEDL
		1				RKLTVDVLPALRANTRALWQV
						EVSAFPSFMTQATPATQPLINAE
						PAVAAPTEQNPQVGPVMPGVQ
		1				GADAPVVAQNGPSRDVKLTFA
						PIAPPPGSMVLRGLNPNGSLEFG
						MRSDEVVTKAMVQLELTPWH
						GYGCQCPFLPVKYLYAAIGFSV
		1				MIEALNQLAIFNRRRFLSANQT
:		1				LRQRTTEAVMRLLSGQKEDAE
				:		LDAETASMLVDHGNQQIFNPQ
		l				ERRMIERVLNLNQRTVSSIMTS
						RHDIEHIDLNAPEEEIRQLLERN
				:		QHTRLVVTDGDDAEDLLGVVH
		l	1	}		VIDLLQQSLRGEPLNLRVLIRQP
		1	1	Ì		LVFPETLPLLPALEQFRNARTHF
		1		-		AFVVDEFGSVEGIVTLSDVTETI
		1				AGNLPNEVEEIDARHDIQKNAD
		1				RSWTGN/GLLPLGDCRL*QASP
		1				TGCINRRGS\HYGYPTLYEFRKR
		1				FLTPGTVCKRCDQRGDAMRVQ
21660	52028	A	21788	558	1563	VQWVFYTYILPLANE TNNRL*K*GNN**LTNQKKSRT
21000	32028	A	21/00	338	1363	1
						RWIHSRILPEAGC/GIASVGRNN CGLWRFTPAKAISWGKIHPVFQ
		1				RQILRVLSQTLFPN/MSLGRNLG
						YGLKMFGVPRAELKPRFKEALP
		İ				MVNLEGFEDRFVDQISGGQQQ
		ł				RVALARALILKPKVLLFDEPLS
						NLDANLRRSMRDKIRELQKQF
						DITSLYVTHDQSEAFAVSDTVL
						VMNKGHIMQIGSPQDLYRQPA
						SRFMASFMGDANLFPATFSDG
						YVDIYGYHLPRPLHFGTQGEG
		1				MVGVRPEAITLSDRGEESQRCV
]						IRHVAYMGPQYEVTVEWHGQE
				1		ILLQVNATRLQPDVGEQYYLEI
		$oxed{oxed}$				HPYGMFVLADAA
21661	52029	Α	21789	201	312	CSWHDRFPDWKAGTHLLARG*
			<u></u>			QLSLPVWLCPVLKLS
21662	52030	Α	21790	925	2388	
21663	52031	Α	21791	1	759	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	ı	in USSN	location of first	codon for last amino acid	,
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21664	52032	Α	21792	1	1086	MEKKITGYTTVDISQWHRKEHF
						EAFQSPAQCTKNQTVQLDITAF
1						LKTVKKNKHKFYPAFIHILARL
ł						MNAHPEFRMAMKGVPGGDSV
						KNTSRVWSGNERDQKLLTENA
						LNDLIPSFLLTGQQTPALGRRVS
						GVIEIADGSRRRKAAALTESDY
]						RVLVGELYDEQMPPLSRLGND
						YRPTSAHPQKRRYASRLQNEFA
						GNISALADAENISRVRDVNLLL
						EVLGVDSVEKTFRVWSGNERD
						QELLMEDALDDLIPSFLLTGQQ
					ļ	TPAFGRRVSGVIEIADGSRRRK
						AAALTESDYRVLVGELDDEQM
						AALSRLGNDYRPTSAYERGQR
						YASRLQNEFAGNISALADAENI
						SRKIIPRCINS\AKLP
21665	52033	A	21793	1	935	MLRDLLKNVDLKGFEPDVRILL
21003	132033		21//3			TKYSNSNGSQSPWMEEQIRDA
						WGSMVLKNVVRETDEVGKGQI
						RMRTVFEQAIDQRSSTDNEGAP
		1				VIPKHTLNTQPVEDTSLSTPAAP
		ŀ				MVDSLIARVGVMARGNAITLP
						VCGRDVKFTLEVLRGDSVEKTS
				]		RVWSGNERDQELLTEDALDDLI
						PSFLLTGQQTPAFGRRVSGVIEI
						ADGGSSRSKKRRRGCEGGRR/V
ļ ·						RQGCPDQKHPAAFSVSGGKIHL
						SGRTPVEKRPRLPQPRDLGARA
						FAACRGSRGRPWNWLPAALRL
						HADLGCVRGGGGGWPGANPY
i i						PAPASSTC
21666	52034	Α	21794	181	923	QHAIVTGDAVGMHHIPQEARQ
21000	32034		21/24	101		YRHNQAYAYSIQGDGAEDDDE
						RIVRFHTRCLTEAFEKLQCGGV
						VNQLSRRGNNQHISSTYDINRA
						DTQVRRAVNNYDIIVMSNSFN\
						ASRSIRCGSVGSLPSSNLPINSVS
						IRCRAR/SG/EE*CQAPASKWAL
			ŀ			
						LHK*HRPFPQDRKAGECLLHEY
						EDLVPIRDTLRLFPGGRYLPRA
						KHVAPSEPDPEQDEQKLRFCKR
						HLYGQQPRSPVEIRLQHVAIAY
		<u> </u>				PTHHAYD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
			<u> </u>			
21667	52035	Α	21795	178	798	IRLEPFKINVLEQITKHIEKLQCG
						GVVKQLSRRGNNQHISSTYDIN
						RADTQVRRAVNNYDIIVMSNSF
						N\ASRSIRCGSVGSLPSSNLPINS
						VSIRCRAR/SG/EE*CQAPASKW
						ALLHK*HRPFPQDRKAGECLLH
						EYEDLVPIRDTLRLFPGGRYLPR
				-		AKHVAPSEPDPEQDEQKLRFCK
						RHLYGQQPRSPVEIRLELVAIA
21668	52036	A	21796	499	972	ҮОТННА
21669	52037	A	21797	3	1141	SRVKLGQSSESQVAALKTREPM
						NIATLEVLVLKGLRVCFVKVNH
						PKGPASMYPGWVPILYIHAEDP
						LLPFYLGEKEDVTYAIKPTCWP
				+		GLDIIPSCLALPRIETELMGKFD
						EGKLPTDPHLMLRLAIETVAHD
	ĺ					YDVIVIDSAPNLGIGTINVVCAA
						DVLIVPTPAELFDYTSALQFFD
				ŀ		MLRPLLKNVDLKGFEPDVRILL
	1					TKYSNSNGSQSPWMEEQIRDA
						WGSMVLKNVKEFAGNISALAD
						AENISRKIITRCINTAKLPKSVV/
	ŀ			1		ALFSHPGELSA\GDALQKAFTD
						KEELLKQQASNLHEQKKAGVIF
	}					EAEEVITLLTSVLKTSSASRTSL
						SSRHQFAPGATVLYKGDKMVL
						NLDRSRVPTECIEKIEAILKELE
21670	52038	Α	21798	1	617	LEPASLSIPNLLLHPRGLRAITIA
	ł					VFGKQNTYIRLEPFKINVLEQIT
	:					KHIEKLQCGGVVKQLSRRGNN
						QHISSTYDINRADTQVRRAVNN
	:					YDIIVMSNSFN\ASRSIRCGSVGS
		1				LPSSNLPINSVSIRCRAR/SG/EE*
						CQAPASKWALLHK*HRPFPQD
						RKAGECLLHEYEDLVPIRDTLR
	<u> </u>	ļ				LFPGGRYLPRAKHVAPSEPILSC
21671	52039	Α	21799	181	953	QHAIVTGDAVGMDDIPQEARQ
						YRHNQAYAYSIQGDGAEDDDE
						RIVRFHTRRLTFKINVLEQITKHI
				1		EKLQCGGVVKQLSRRGNNQHI
						SSTYDINRADTQVRRAVNNYDI
						IVMSNSFN\ASRSIRCGSVGSLPS
						SNLPINSVSIRCRAR/SG/EE*CQ
						APASKWALLHK*HRPFPQDRK
						AGECLLHEYEDLVPIRDTLRLFP
						GGRYLPRAKHVAPSEPDPEQDE
						QKLRFCKRHLYGQQPRSPVEIR
		<u> </u>	ļ			LQHVAIAYQTHHAYD
21672	52040	A	21800	1 .	798	
21673	52041	A	21801	1	1782	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible πucleofide insertion)
21674	52042	Α	21802	666	1131	KGRGTYGNERDQELLT\EDALD
1						DLIPSFLLTGQQTPAFGRRVSGV
į						IEIADGSRRRKAAALTESDYRV
1		ŀ			}	LVGELDDEQMAALSRLGNDYR
						PTSAYERGQRYASRLQNEFAG
Ì		ľ				NISALADAENISRKIITRCINTRO
		ļ				LPKSVVGFFSHPGELSARSGDA
21675	52043	Α	21803	1	3297	
21676	52044	Α	21804	1	898	
21677	52045	Α	21805	1	1977	
21678	52046	Α	21806	1	1929	
21679	52047	Α	21807	181	755	QHAIVTGDAVGMDDIPQEARQ
						YRHNQAYAYSIQGDGA\GMTM
-						SALLDFIHGA***YIRCGSVGRL
						PSSNLPINSVSIRCRAR/SG/EE*C
			:			QAPASKWALLHK*HRPFPQDR
						KAGECLLHEYEDLVPIRDTLFL
İ						FPGGRYLPRAKHVAPSEPDPEQ
						DEQKLRFCKRHLYGQQPRSPVE
						IRLQHVAIAYQTHHAYD
21680	52048	Α	21808	181	456	QHAIVTGDAVGMDDIPQEARQ
						YRHNQAYAYSIQGDGAEDDDE
ļ						RIVRFHTRCLA*SRRAVNNYDII
		i				VMSNSFNGQSEHQVWIGGQFT
<u> </u>						FIKFAH
21681	52049	Α	21809	1	780	
21682	52050	Α	21810	1	1050	
21683	52051	Α	21811	5	735	PPAAP/MGNLLMARVGVKARG
						KAHYLPLLRGESVRKTSRVGSG
		ļ				NERDQELLTEDALDDLIPSFLLT
						GQQTPAFGRRVSGVIEIADGSR
					İ	RRKAAALTESDYRVLVGELDD
ŀ					†	EQMAALSRLGNDYRPTSAYER
		ŀ			1	GQRYASRLQNEFAGNISALADA
						ENISRQIITRCINTAKLPKSVVAL
						FSHPGELSARSGDALQKAFTEK
						RTLKQQALTSEQKKRGI/YEAE
	-		<u> </u>	<u> </u>		EVSSSLSPQGRLSNRSASKDMY

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21684	52052	A	21812	1	887	MNMNIKKIVKQATVLTFTTALL
1		ļ				AGGATQAFAKENNQKAYKETK
		1				AVGSGNERDQELLTEEAWNDL
		ļ				ILSFLLTGQQTPAFGRRVSGVIEI
		l				PDGSRRRKAAALPKSDYRVLV
						GELDDEQMAALSRLGNDYRPT
						SAYERGQRYASRLQNEFAGNIS
						ALADAENISRKIITRCINTAKLP
						KSVVALFSHPGELSARSGDALQ
				1		KAFTDKEELLKQQASNLHEQK
				}		KAGVIFEAEEVITLLTSVLKTSS
				1		ASRTSLSSRHQFAPGATVLYKG
						DKMVLNLDRSRVPTEC/IEKIEA
						ILKELEKPAP
21685	52053	A	21813	345	921	RWRLQNLSFCSSCSGSGSEGAT
		ļ				CFARGSAYELGQRYASRLQNA
						FAGKISALADAEIFSRKII\TRCIN
						TAKLLKSVVALFSHPVELSARS
				1		GDALQKAFTDKEELLKQQASN
						LHEQKKAGVIFEAEEVITLFTSV
						LKTSSASRTSLSSRHQFAPGAT
						VLYKGDKMVLNLDRSRVPTEC
						IEKIEAILKELEKPAP
21686	52054	Α	21814	1	1989	
21687	52055	A	21815	6	311	
21688	52056	Α	21816	3	848	PGGDSVKNTSRVWSGNERDQK
		1				LLTENALNDLIPSFLLTGQQTPA
				1		LGRRVSGVIEIADGSRRRKAAA
İ						LTESDYRVLVGELYDEQMPPLS
						RLGNDYRPTSAHPQKRRYASRL
				1		QNEFAGNISALADAENISRVRD
						VNLLLEVLGVDSVEKTFRVWS
				-		GNERDQELLMEDALDDLIPSFL
						LTGQQTPAFGRRVSGVIEIADG
						SRRRKAAALTESDYRVLVGEL
				1		DDEQMAALSRLGNDYRPTSAY
						ERGQRYASRLQNEFAGNISALA
01555		<u> </u>	0.015	ļ. ———	0.60	DAENISRKIIPRCINS\AKLP
21689	52057	A_	21817	1	969	
21690	52058	Α	21818	1	1614	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			}	sequence		
21601	52050	1	121010	<u> </u>	1022	MDDIDOEADOVRIDIOAVAVOL
21691	52059	A	21819	1	832	MDDIPQEARQYRHNQAYAYSI
]		1				QGDGAEDDDERIVRFHTRVTV
						DSDTLASDAARLTCRHGLARD
						VKFTLEVLRGDSVEKTSRVWS
						GNERDQELLTEDALDDLIPFFLL
		1				TGQQTPAFGRRVSGVIEIADGS
		ĺ				RRRKAAALTESDYRVLVGELD
						DEQMAALSRLGNDYRPTSAYE
						RGQRYASRLQNEFAGNISALAD
						AENISRKIITR\CINT\AQLAKSG
	i					VALFSHPGELSAPSGDAL*KAF
		l	:			TD*EEFL*PQGSNLY\EQKKAG\
						VIFGS*KKFSLFLTSVV
21692	52060	Α	21820	678	1338	TLRQTKPDNSAEIADGSRRRKA
						AALTE\SDYRVLVGELDDKQM
		İ				AALSRLGNDYRPTSAYERGQR
						YASRLQNEFAGNISALADAENI
						SRKIITRCINTAKLPKSVVALFS
						HPGELSARSGDALQKAFTDKEE
						LLKQQASNLHEQKKAGVIFEAE
						EVITLLTSVLKTSSASRTSLSSR
						HQFAPGATVLYKGDKMVLNLD
						RSRVPTECIEKIEAILKELEKPAP
21693	52061	Α	21821	I	1809	
21694	52062	Α	21822	2	1010	
21695	52063	Α	21823	284	2177	
21696	52064	Α	21824	1	1296	
21697	52065	В	21825	1	1044	
21698	52066	Α	21826	637	1073	IRLEPFKINVLEQITKHIEKLQCG
						GVVKQLSRRGNNQHISSTYDIN
						RADTQVRRAVNNYDIIVMSNSF
						N\ASRSIRCGSVGSLPSSNLPINS
						VSIRCRAR/SG/EE*CQAPASKW
						ALLHK*HRPFPQDRKAGECLLH
						EYEDLYPSVIH
21699	52067	Α	21827	1	2403	
	1			L	L	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
	İ			sequence		
21700	52068	Α	21828	1471	3186	LPHPDMEIRGRVEQRVGYTIEQI
						NHMRDVFGTRLRRAEDVFTPVI
						GVAAIKTGTTVHIVSSSSLLGG
						DHGPSRIGGLLLVCDYGPLVSS
						SGISGPTRIVGLIISLGPGYHLCF
						RSDYRTGITDPLYIGDATIRVSL
						RVYSLPSRSRIEHVVANCGMIP
						KQQQNEKYQVPQFDQSTIKNIE
			į			SAKGLDVWDSWPLQNADGTV
	İ			+		AEYNGYHVVFALAGSPKDADD
						TSIYMFYQKVGDNSIDSWKNA
						GRVFKDSDKFDANDPILKDQTQ
						EWSGSATFTSDGKIRLFYTDYS
						GVMARGNAITLPVCGRDVKYT
						LEVLRGDSVEKTSRVWSGNER
						DQELLTEDALDDLIPSFLLTGQ
						QTPAFGRRVSGVIEIADGSRRR
						KAAALTESDYRVLVGELDDEQ
						MAALSRLGNDYRPTSAYERGQ
						RYASRLQNEFAGNISALADAEN
						ISRKIITRCINTAKLPKSVVALFS
						HPGELSARSGDALQKAFTDKEE
						LLMQQASNLHEQKKAGVIFEA
						EEVITLLTSVAKTSSASRTSLSS
						RHQF\APGATV\LYKGDKMVLN
						LDMSR\VQLECLDRLHACRSTL
21701	52069	Α	21829	158	926	AGPVIQNYVNTQPV/EETSYSTP
						AAPMVDSLIARVGVMAPGNAI
						TLPVCGRDVKFTLEVLRGDSVK
						KTSRVWSGNERDQELLTEDAL
						DDLIPSFLLTGQQTPAFGRRVSG
						VIEIADGSRRRKAAALTESDYR
						VLVGELDDEQMAALSRLGNDY
						RPTSAYERGQRYASRLQNEFAG
						NISALADAENISRKIITRCINTAK
						LPKSVVALFSHPGELSARSGDA
						LQKAFTDKEELLKQQGGSSRV
						GDNSIDSWKTRGRVL
21702	52070	В	21830	1	3774	
21703	52071	В	21831	1	1495	
21704	52072	Α	21832	556	1622	
21705	52073	Α	21833	1	2616	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	4	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21706	52074	Α	21834	1412	3310	SVTSCLKVTSVCLTSAAASCRS
						VLMRSTAELQCASHTLRKRKA
						ARRLISLMMTARKARKITRRW
						RIGEAADLVGVSSQAIRDAEKA
						GRLPHPDMEIRGRVEQRVGYTI
						EQINHMRDVFGTRLRRAEDVFP
						PVIGVAAHKGNDPQGTASMYH
						GWVPDLHIHAEDTLLPFYLGEK
		l				DDVTYAIKPTCWPGLDIIPSCLA
		İ				LHRIETELMGKFDEAQPNLGIG
						TINVVCAADVLIVPTPAELFDY
}						TSALQFFDMLRDLLKNVDLKG
						FEPDVRILLTKYSNSNGSQSPW
j						MEEQIRDAWGSMVLKNVVRET
						DEVGKGQIRMRTVFEQAIDQRS
						STDTSLSTPAAPMVDSLIARVG
						VMARGNAITLPVCGRDVKFTL
		1				EVLRGDSVEKTSRVWSGNERD
						QELLTEDALDDLIPSFLLTGQQT
						PAFGRRVSGVIEIADGSRRRKA
						AALPE\SNYGVLVAELDDEQMP
		ŀ				AYPRLGNDYRPTSAYERGQRY
				:		ATRLQNEFAG\NIS\ALADAENIS
		l .				\IITRCINTAKLPKSVVALFSHPG
						ELSARSGDALQKAFTDKEELLK
						QQASNLHEQKKAGVIFEAEEVI
						TLLTSVLKTSSASRTSLSSRHQF
						APGATVLYKGDKMVLNLDRSR
						VPTECIEKIEAILKELEKPAP
21707	52075	Α	21835	783	2371	
21708	52076	Α	21836	3236	4718	
21709	52077	Α	21837	1	2481	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence ( X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21710	52078	A	21838	2450	3963	LLTAPITGGIGTINVVCADDVMI VPTPAELFDYTSALQFFDMLRD LLKNVDLKGFEPDVRILLTKYS NSNGSQSPWMEEQIRDAWGSM VLKNVVRETDEVGKAAPMVDS LIARVGVMARGNAITLPVCGRD VKFTLEVLRGDSVEKTSRVWS GNERDQELLTEDALDDLIPSFLL TGQQTPAFGQR\VSGVKEIADGI RRRKAAALTE\SDYRVLVGELD DEQMAALSRLGNDYRPTSAYE RGQRYASRLQNEFAGNISALAD AENISRKIITRCINTAKLPKSVV ALFSHPGELSARSGDALQKAFT DK\EDLLKQQASNLHGAEKSW GCYFEAEEVITLLTSVLKTS\A SRTSLSSRHQFAPGATVLYKGD KMVLNLDRSRVPTEYRYGVFL NVLLPATKYFQLHDPSLRNMEL NYSLASEDLVLKGLRVLLVEG NDPQGTASMYHGWVPDLHIHA EDTLLPFYLGEKDDVTYAIKPT CWPGLDIIPSCLALHRIETEYNG
21711	52079	A	21839	1573	2858	EPVFEQAIDQRSSTDTSLSTPAA PMVDSLIARVGVMARGNAITLP VCGRDVKFTLEVLRGDSVEKTS RVWSGNERDQELLTEDALDDLI PSFLLTGQQTPAFGRRVSGVIEI ADGSRRRKAAALTESDYRVLV GELDDEQMAALSRLGNDYRPT SAYERGQRYASRLQNEFAGNIS ALADAENISQ*ICWKYFCAG*C GKYF\RKIITRCINTAKLPKSVV ALFSHPGELSARSGDALQKAFT DKEELLKQQASNLHEQKKAGV ISPPEEVITLLTSEIKTSSASRTSL SSRHQFAPGATVLYKGDKMFIT VKIAKRSQAPCMKSNNALIVIL GTVTLDAVGIGLVMPVLPGLLR DIVHSDSIASHYGVLLALYALM QFLCAPVLGALSDRFGRRPVLL ASLLGATIDYAIMATTPVLWIY PLVNSPSC
21712	52080	В	21840	63	368	
21713	52081	A	21841	189	204	SLPSGIKPQSSSKASSRIQVAPT M*SLPRQPLAWPSGHEACPQG
21714	52082	Α	21842	153	370	LIAGAFRKASGGNNIFRRPKTL* LRRQPR*PQKSTPRRNKLGHYA IIKFPLTTESAVKKIEETNTLVFT VLV

RRNWLRNAEPP KRQGRATKPK RVRDVALRPYS KIRMSLTFRRP PRKSTPRRNKL ESAVKKIEENN IKHQIRQAVKK
RRNWLRNAEPP KRQGRATKPK RVRDVALRPYS KIRMSLTFRRP PRKSTPRRNKL ESAVKKIEENN
KRQGRATKPK RVRDVALRPYS KIRMSLTFRRP PRKSTPRRNKL ESAVKKIEENN
KRQGRATKPK RVRDVALRPYS KIRMSLTFRRP PRKSTPRRNKL ESAVKKIEENN
RVRDVALRPYS KIRMSLTFRRP PRKSTPRRNKL ESAVKKIEENN
KIRMSLTFRRP PRKSTPRRNKL ESAVKKIEENN
PRKSTPRRNKL ESAVKKIEENN
ESAVKKIEENN
≀KHQIRQAVKK
EERRHDEQNNIL
TRNRARTETNT
LISFHSYYSDTS
SGQCTYDRLG
LQHTYDIQSTQ
/NLRLYCANYG
ELLILTRSSNSK
AHPKAEAKAKA
VRSHT/RKKKIR
RLRRQPRYPRK
AIIKFPLTT
VALSLRPGFMS
RRSVTQASGAS
DKTAPKAKEAP
ALKPKKAVVER
MSPTFRRPKTL
STPRRNKLGHY
AGKKNKKNNTP
HQI\RQAVKK\L
LICPDKEKKAY
OVANKIGII
YRIFAVL*SVQ*
GELP*SVAIQPN
/LPFGKCGSTY
1
VKVCSVSVKLL
EGLDRLSERSG
ALDAASOLVSL I
ALDAASQLVSL VSSL/SSTYV*SP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21723	52091	A	21851	11	766	SGRKEANY\YGSLTQAGTVSLG
21,723	2207.	ľ.				LDAEGQEVFVPFSAVLPMVAP
	ł					NDLVFDGWDISSLNLAEAMRR
					-	AKVLDWGLQEQLWPHMEALR
ŀ						P/LGLAVYIPEFIAANQSARA\D
	ė					N\LIPGSRAQQLEQIRRDIRDFRS
						SAGLDKVIVLWTANTERFCEVI
						PGLNDTAENLLRTIELGLEVSP\
						STLF\AVASILGGLCLSFNGSPQ
						NTLVPGALELAWQHRILVGGD
						DFKSGQTKVKSVLVDFLIGSGL
						KTMSIVSYNHLGNNDGEN
21724	52092	A	21852	2	2122	WKSWMVFTAMKLVKAHHRLK
21724	32072	`	21032	2		LRLKGYLPVTCLEGTRNQRIAA
						TIRHNRARGRHQITAMSEIVRE
						LSQLGWDDNKIGKELGMDSDE
						VLRLKQINGLQELFADRQYSRA
	-					WTLKLAQAFSCCQQRRFLLGK
			1			TETNQFMWEGIGAKHRQRNRR
						YAVLLRQPLAKLNIIQARYPNQ
						HRQHAGECNPHCQAGGDIVDR
	1					CPARCFTKGDSKDGSPPQATWP
						FHNSATMTAKRLSEASQCGSK
ļ						KSILRERIIYRPSITRYPFQSTMN
						HPLPEQTLGCIATDYGSSPSPDR
		l				SPGPAHYYWTDHSTANIRYQV
ł						AAEENGEYTYVERVKIPLDHGT
		1				LLIMERATQADWQLEQIRRDIR
1						DFRSSAGLDKVIVLWTANTERF
						CEVIPGLNDTAENLLRTI/D/RSV
						WEVSPSTLFAVASILEGCAFLN
			:			GSPQNTLVPGALELAWQHRVF
						VGGDDFKSGQTKVKSVLVDFLI
						GFRLQRP/SSIVSYNHLG\NNDG
						ENLSAPLQFRSKEV\SRSNVVD
		1				DMVA\SNP\VLYTPGEEPDHCV
Ì						VIKYVPYVGDS*RAL\DEYTSEL
		1			]	MLGGTNTLVL\HNTC\EDSLL\A
						APIML\DLGAA*PSCCQRVTFLA
					1	LDIGTPSRRPFHPVLVPAQAFFF
						KGATLSPPGKP/VLVNAAFSAS
						AAAIENI\LRACVGLPPQNHMLL
						EHKMERPGPSLKRVWTPWLPP
i						YPMLNKKGPGSRCQPMAAPVD
21725	52093	A	21853	1	1908	The state of the s
21723	52093	B	21854	1	1833	
21/20	32034	Пр	121034		11033	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21727	52095	Α	21855	1	749	MFQLMKASSGGEAACMPAGL
						MENYKHTTVLLDEAVNGLNIR
l						PDGIYIDGTFGRG\LGEEGRLLA
						IDRDPQAIAV/AKTIDDPRFSIIH
		l				GPFSALGEYVAERDLIGKIDGIL
						LDLGVSSPQLDDAERGFSFMRD
						GPLDMRMDPTRGQSAAEWLQT
						AEEADIAWVLKTYGEERFAKRI
						ARAIVERNREQPNDPELEEIEQA
						LKSSLNVLAPGGGVSIISFHSLE
						DRIVK/RFMREHQFCVVVRSGV
						GLKT*SVR
21728	52096	A	21856	532	1418	CFASGFIGELRKNIVGFIEQDHT
						RGLVFRLTPGRTLRAESLNLVIC
		1				LPMNFQPYDSDADWVITGVPF
						DMATSGRAGGRHGPAAIRQVS
						TNLAWEHNRFPWNFDMRERLN
						VVDCGDLVYAFGDAREMSEKL
						QAHAEKLLAAGKRMLSFGGDH
						FVTLPLLRAHAKHFGKMALVH
						FDAHTDTYANGCEFDHGTMFY
						TAPKEGLIDPNHSVQIGIRTEFD
						KDNGFTVLDACQVNDRSVDDV
						IAQVKQIVGDMP/VYLTLYNCG
						GPS*QFHSLRPAQSPLDRCRAR
		<u> </u>				YLAPQCVPDFRTLQR
21729	52097	В	21857	64	1842	
21730	52098	Α	21858	1	503	MRWLVSPIEDTGHKALLFACL
						ALHRACVSGHCLSPKCVGVDG
				}		RSHEMVAPGLFLDWLLDGTGV
				1		WDGTSTNVEGAQEHAVLILLII
						TDGVISDMEETGMPWCRLPSCP
						CPSSSGRGQCGLRCHGVPGWG
						RRMLRSHTGEEAARDIVSSFPF
						ESSATMSEEVYDPRSLYERLQE
						QRTGSSRTARQEAVVGSKVTV
						SQPGHRQETTVLEHLVQY/FILL
						IITDGVISDMEE/TRHAVVQASK
						LPMSIII/VGVGNADFAAMEFLD
				1		GDA/GMLRSHTGEEAARDI/VQ
				_		FVPFREFRNNVRGGL*PSISI
21731	52099	Α	21859	2223	2363	

Sequence   09/540,217   codon for peptide sequence   deteition, "possible nucleotide insertion sequence   Sequence   deteition, "possible nucleotide insertion   Sequence   Se	SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
21732   52100   A   21860   3   693   GDDGKWMLVHRTEVIKYTLE	NO:	1	hod		1		*=Stop codon, /=possible nucleotide
21732   52100   A   21860   3   693   GDDGKWMLVHRTEVIKYTLL		sequence		09/540,217	1 ' '	of peptide sequence	deletion, \=possible nucleotide insertion)
VWKPFTVPLVSLCDGDMEKP VMCYDYDNDGGHDFIGEFGTT MSQMCARDSVPLEFECINP KQRKKNYKNSGIIILRSCKIN DYSFLDYLLGGGQLIHVSPLGI FTASNGGPLDPSSLHYINPM TNEYLSAIWAVGQIIQDYDSD MFPALGFGAQLPPDWKVSHE AINFNPTPFCSGVDGIAQAIY TCLPHIRFYGFT  121733 \$2101 A 21861 1 2231 MYSQEONGFCNRTDCQELLP PAPMAHIPSGGAPAGAAPM PQYCVCKVELSVSGQNLLDRI VTSKSDPFCVLFTENNGRWIE DRTETAINNLNPAFSKFVLD HFEEVGKLKFALFDQDKSSMI LDEHDFLQGFSCSLGTIVSSK TRPLLLLNDKPAGKGLITIAA LSDNRVITI.SLACRRLDKKDL GKSDPFLEFYKPGDDGKWML HRTEVVIKYTI.DPVWKPFTVF VSLCDGDMEKPIQVMCYDVT DGGHDFIGEFQTSVFKKGEA DSVPLEFECINFKKQRK.KNY NSGIIILRSCKVNQRGQABARK QVSGHTEGTDRALDLEPARP KSRLCHILALWPSELHLTRQ HLWNGDINEAEHVQGYSSLT CGGGGRHELTFQGLSWENG CCELLPPTVGMQGAEDIGWGH VLSGISVTVTFMINRDYSFLD LGGCQUMFTVGIDFTASNGN DPSSLHYINPMGTNEYLSAIW VGQIIQDYDRRSFLKPDFWA TEAVTGQHWFSVGTTASNGN DPSSLHYINPMGTNEYLSAIW VGQIQDYDRRSFLKPDFWA TEAVTGQHWFSVGTTASNGN QAYSACLPHIRFYCPTTSPIV HVARFAAQATQQRTATQYFII IITDGVISDMEETRHAVVQASI LPMSIIIVGVGNADAFAMFELI GDSPHAALPTRGRRQPRDIVQ QAYSACLPHIRFYCPTFSPIV HVARFAAQATQQRTATQYFII IITDGVISDMEETRHAVVQASI LPMSIIIVGVGNADAFAMFELI GDSPHAALPTRGRRQPRDIVQ QOSLSNIREAGAVPRFFARRYV GRIUG*FLFLGVLRGAMPFFARRYV FTAKAEAGDRTVA*PHKPKM		[			sequence		
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LDEHDFLGQFSCSLGTIVSSKK TRPLLLLNDKPAGKGLITIAAG LSDNRVITLSLACRRLDKKDL GKSDPFLEFYKPGDDGKWML HRTEVVIKYTLDPVWKPFTVF VSLCDGDMEKPIQVMCYDYD DGGHDFIGEFQTSVFVKMCEA: DSVPLEFECINPKQRKKKNY NSGIIILRSCKVNQRGQARARG QVSGHTEGTDRALDLEPARPY KSRLCHILALWPSELHLTRPQ: HLWNGDINEAEHVQGYSSLTI CGGQGRHELTFQGLSWWENG CELLPPTVGMQGAEDIGWGH VLSGISVTVTPMINRDYSFLDY LGGCQLMFTVGIDFTASNGNE DPSSLHYINPMGTNEYLSAIW VGQIIQDYDRPRSFLKPDPWA TEAVTGQHVMFSVYETHVEW RQIGLPDVANQNTRHPVKFEF VSHEFAINFNPTNPFCSGVDGI QAYSACLPHIRFYGPTNFSPIV HVARFAAQATQQRTATQYFII IITDGVISDMEETRHAVVQASI LPMSIIIVGVGNADFAAMEFLI GDSPHAALPTRGRRQPRDIVQ 21734 52102 A 21862 694 825 QPGLSINIEAGAVPRFFARRYV GNLG*FLFLGVLRCFSSPGSPE 21735 52103 A 21863 1032 1127 FYAKAEAGDRTVA*PHKPKM							DRTETAINNLNPAFSKKFVLDY
TRPLLLLNDKPAGKGLITIAAG LSDNRVITLSLACRRLDKKDL GKSDPFLEFYKPGDDGKWML HRTEVVIKYTLDPVWKPFTVF VSLCDGDMEKPIQVMCYDYD DGGHDFIGEFQTSVFVKMCEA DSVPLEFECINPKKQRKKKNY NSGIIILRSCKVNQRGQARARG QVSGHTEGTDRALDLEPARPV KSRLCHILALWPSELHLTRPQ HLWNGDINEAEHVQGYSSLTI CGGQGRHELTFQGLSWWENG CELLPPTVGMQGAEDIGWGH VLSGISVTVTPMINRDYSFLDV LGGCQLMFTVGIDFTASNGNI DPSSLHYINPMGTNEYLSAIW VGQIIQDYDRPRSFLKPDPWA TEAVTGQHVWFSVYETHVEW RQIGLPDVANQNTRHPVKFEF VSHEFAINFNPTNPFCSGVDGI QAYSACLPHIRFYGPTNFSPIV HVARFAAQATQQRTATQVFII IITDGVISDMEETRHA VVQASI LPMSIIIVGVGNADFAAMEFLI GDSPHAALPTRGRRQPRDIVQ 21734 52102 A 21862 694 825 QPGLSINIEAGAVPRFFARRYV GNLG*FLFLGVLRCFSSPGSPH 21735 52103 A 21863 1032 1127 FYAKAEAGDRTVA*PHKPKM							HFEEVQKLKFALFDQDKSSMR
LSDNRVITLSLACRRLDKKDL GKSDPFLEFYKPGDDGKWML HRTEVVIKYTLDPVWKPFTVF VSLCDGDMEKPIQVMCYDYD DGGHDFIGEFQTSVF\KMCEA: DSVPLEFECINPKQRKKKNY NSGIIILRSCKVNQRGQARARG QVSGHTEGTDRALDLEPARP\ KSRLCHILALWPSELHLTRPQ HLWNGDINEAEHVQGYSSLT\ CGGQGRHELTFQGLSWWENG CELLPPTVGMQGAEDIGWGH VLSGISVTVTPMINRDYSFLD\ LGGCQLMFTVGIDFTASNGNI DPSSLHYINPMGTNEYLSAIW VGQIIQDYDRPRSFLKPDPWA TEAVTGQHVMFSVYETHVEW RQIGLPDVANQNTRHPVKFEF VSHEFAINFNPTNPFCSGVDGI QAYSACLPHIRFYGPTNFSPIV HVARFAAQATQQRTATQYFII IITDGVISDMEETRHAVVQASI LPMSIIIVGVGNADFAAMEFLI GDSPHAALPTRGRRQPRDIVQ 21734 52102 A 21862 694 825 QPGLSINIEAGAVPRPFARRYV GNLG*FLFLGVLRCFSSPGSPH 21735 52103 A 21863 1032 1127 FYAKAEAGDRTVA*PHKPKM			İ				LDEHDFLGQFSCSLGTIVSSKKI
GKSDPFLEFYKPGDDGKWML HRTEVVIKYTLDPVWKPFTVF VSLCDGDMEKPIQVMCYDYD DGGHDFIGEFQTSVFKMCEA. DSVPLEFECINPKKQRKKKNY NSGIIILRSCKVNQRGQARARG QVSGHTEGTDRALDLEPARPV KSRLCHILALWPSELHLTRPQ: HLWNGDINEAEHVQGYSSLTI CGGQGRHELTFQGLSWWENG CELLPPTVGMQGAEDIGWGH VLSGISVTVTPMINRDYSFLDL LGGCQLMFTVGIDFTASNGNI DPSSLHYINPMGTNEYLSAIW VGQIIQDYDRPRSFLKPDPWA TEAVTGQHVMFSVYETHVEW RQIGLPDVANQNTRHPVKFEF VSHEFAINFNPTNPFCSGVDGI QAYSACLPHIRFYGPTNFSPIV HVARFAAQATQQRTATQYFII IITDGVISDMEETRHAVVQASI LPMSIIIVGVGNADFAAMEFLI GDSPHAALPTRGRRQPRDIVQ 21734 52102 A 21862 694 825 QPGLSINIEAGAVPRPFARRYV GNLG*FLFLGVLRCFSSPGSPH 21735 52103 A 21863 1032 1127 FYAKAEAGDRTVA*PHKPKM							TRPLLLLNDKPAGKGLITIAAQE
HRTEVVIKYTLDPVWKPFTVE VSLCDGDMEKPIQVMCYDYD DGGHDFIGEFQTSVF\KMCEA: DSVPLEFECINPKKQRKKKNY NSGIIILRSCKVNQRGQARARG QVSGHTEGTDRALDLEPARPV KSRLCHILALWPSELHLTRPQ! HLWNGDINEAEHVQGYSSLTI CGGQGRHELTFQGLSWWENG CELLPPTVGMQGAEDIGWGH VLSGISVTVTPMINRDYSFLDY LGGCQLMFTVGIDFTASNGNF DPSSLHYINPMGTNEYLSAIW VGQIIQDYDRPRSFLKPDPWA TEAVTGQHVMFSVYETHVEW RQIGLPDVANQNTRHPVKFEF VSHEFAINFNPTNPFCSGVDGI QAYSACLPHIRFYGPTNFSPIV HVARFAAQATQQRTATQYFII IITDGVISDMEETRHAVVQASI LPMSIIIVGVGNADFAAMEFLI GDSPHAALPTRGRRQPRDIVQ  21734 52102 A 21862 694 825 QPGLSINIEAGAVPRPFARRYV GNLG*FLFLGVLRCFSSPGSPH 21735 52103 A 21863 1032 1127 FYAKAEAGDRTVA*PHKPKM			1				LSDNRVITLSLACRRLDKKDLF
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NSGIIILRSCKVNQRGQARARG QVSGHTEGTDRALDLEPARPV KSRLCHILALWPSELHLTRPQI HLWNGDINEAEHVQGYSSLTI CGGQGRHELTFQGLSWWENG CELLPPTVGMQGAEDIGWGH VLSGISVTVTPMINRDYSFLDY LGGCQLMFTVGIDFTASNGNF DPSSLHYINPMGTNEYLSAIW VGQIIQDYDRPRSFLKPDPWA TEAVTGQHVMFSVYETHVEW RQIGLPDVANQNTRHPVKFEF VSHEFAINFNPTNPFCSGVDGI QAYSACLPHIRFYGPTNFSPIV HVARFAAQATQQRTATQYFII IITDGVISDMEETRHAVVQASI LPMSIIIVGVGNADFAAMEFLI GDSPHAALPTRGRRQPRDIVQ 21734 52102 A 21862 694 825 QPGLSINIEAGAVPRPFARRYV GNLG*FLFLGVLRCFSSPGSPH 21735 52103 A 21863 1032 1127 FYAKAEAGDRTVA*PHKPKM							DGGHDFIGEFQTSVF\KMCEAR
QVSGHTEGTDRALDLEPARPY KSRLCHILALWPSELHLTRPQ HLWNGDINEAEHVQGYSSLTI CGGQGRHELTFQGLSWWENG CELLPPTVGMQGAEDIGWGH VLSGISVTVTPMINRDYSFLDY LGGCQLMFTVGIDFTASNGNF DPSSLHYINPMGTNEYLSAIW VGQIIQDYDRPRSFLKPDPWA TEAVTGQHVMFSVYETHVEW RQIGLPDVANQNTRHPVKFEF VSHEFAINFNPTNPFCSGVDGI QAYSACLPHIRFYGPTNFSPIV HVARFAAQATQQRTATQYFII IITDGVISDMEETRHAVVQASI LPMSIIIVGVGNADFAAMEFLI GDSPHAALPTRGRRQPRDIVQ 21734 52102 A 21862 694 825 QPGLSINIEAGAVPRPFARRYV GNLG*FLFLGVLRCFSSPGSPF 21735 52103 A 21863 1032 1127 FYAKAEAGDRTVA*PHKPKM					1		DSVPLEFECINPKKQRKKKNYK
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HLWNGDINEAEHVQGYSSLTI CGGQGRHELTFQGLSWWENG CELLPPTVGMQGAEDIGWGH VLSGISVTVTPMINRDYSFLDY LGGCQLMFTVGIDFTASNGNF DPSSLHYINPMGTNEYLSAIW VGQIIQDYDRPRSFLKPDPWA TEAVTGQHVMFSVYETHVEW RQIGLPDVANQNTRHPVKFEF VSHEFAINFNPTNPFCSGVDGI QAYSACLPHIRFYGPTNFSPIV HVARFAAQATQQRTATQYFII IITDGVISDMEETRHAVVQASI LPMSIIIVGVGNADFAAMEFLI GDSPHAALPTRGRRQPRDIVQ C1734 52102 A 21862 694 825 QPGLSINIEAGAVPRPFARRYV GNLG*FLFLGVLRCFSSPGSPH C1735 52103 A 21863 1032 1127 FYAKAEAGDRTVA*PHKPKM							QVSGHTEGTDRALDLEPARPVF
CGGQGRHELTFQGLSWWENG CELLPPTVGMQGAEDIGWGH VLSGISVTVTPMINRDYSFLDY LGGCQLMFTVGIDFTASNGNF DPSSLHYINPMGTNEYLSAIW VGQIIQDYDRPRSFLKPDPWA TEAVTGQHVMFSVYETHVEW RQIGLPDVANQNTRHPVKFEF VSHEFAINFNPTNPFCSGVDGI QAYSACLPHIRFYGPTNFSPIV HVARFAAQATQQRTATQYFII IITDGVISDMEETRHAVVQASI LPMSIIIVGVGNADFAAMEFLI GDSPHAALPTRGRRQPRDIVQ 21734 52102 A 21862 694 825 QPGLSINIEAGAVPRPFARRYV GNLG*FLFLGVLRCFSSPGSPH 21735 52103 A 21863 1032 1127 FYAKAEAGDRTVA*PHKPKM							KSRLCHILALWPSELHLTRPQSP
CELLPPTVGMQGAEDIGWGH VLSGISVTVTPMINRDYSFLDV LGGCQLMFTVGIDFTASNGNF DPSSLHYINPMGTNEYLSAIW VGQIIQDYDRPRSFLKPDPWA TEAVTGQHVMFSVYETHVEW RQIGLPDVANQNTRHPVKFEF VSHEFAINFNPTNPFCSGVDGI QAYSACLPHIRFYGPTNFSPIV HVARFAAQATQQRTATQYFII IITDGVISDMEETRHAVVQASI LPMSIIIVGVGNADFAAMEFLI GDSPHAALPTRGRRQPRDIVQ CONTRACTOR OF THE STATE OF T							HLWNGDINEAEHVQGYSSLTFL
VLSGISVTVTPMINRDYSFLDY LGGCQLMFTVGIDFTASNGNF DPSSLHYINPMGTNEYLSAIW VGQIIQDYDRPRSFLKPDPWA TEAVTGQHVMFSVYETHVEW RQIGLPDVANQNTRHPVKFEF VSHEFAINFNPTNPFCSGVDGI QAYSACLPHIRFYGPTNFSPIV HVARFAAQATQQRTATQYFII IITDGVISDMEETRHAVVQASI LPMSIIIVGVGNADFAAMEFLI GDSPHAALPTRGRRQPRDIVQ 21734 52102 A 21862 694 825 QPGLSINIEAGAVPRPFARRYV GNLG*FLFLGVLRCFSSPGSPH 21735 52103 A 21863 1032 1127 FYAKAEAGDRTVA*PHKPKM							CGGQGRHELTFQGLSWWENCH
LGGCQLMFTVGIDFTASNGNF DPSSLHYINPMGTNEYLSAIW VGQIIQDYDRPRSFLKPDPWA TEAVTGQHVMFSVYETHVEW RQIGLPDVANQNTRHPVKFEF VSHEFAINFNPTNPFCSGVDGI QAYSACLPHIRFYGPTNFSPIV HVARFAAQATQQRTATQYFII IITDGVISDMEETRHAVVQASI LPMSIIIVGVGNADFAAMEFLI GDSPHAALPTRGRRQPRDIVQ  21734 52102 A 21862 694 825 QPGLSINIEAGAVPRPFARRYV GNLG*FLFLGVLRCFSSPGSPH 21735 52103 A 21863 1032 1127 FYAKAEAGDRTVA*PHKPKM							CELLPPTVGMQGAEDIGWGHP
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VGQIIQDYDRPRSFLKPDPWA TEAVTGQHVMFSVYETHVEW RQIGLPDVANQNTRHPVKFEF VSHEFAINFNPTNPFCSGVDGI QAYSACLPHIRFYGPTNFSPIV HVARFAAQATQQRTATQYFII IITDGVISDMEETRHAVVQASI LPMSIIIVGVGNADFAAMEFLI GDSPHAALPTRGRRQPRDIVQ  21734 52102 A 21862 694 825 QPGLSINIEAGAVPRPFARRYV GNLG*FLFLGVLRCFSSPGSPH  21735 52103 A 21863 1032 1127 FYAKAEAGDRTVA*PHKPKM			1				LGGCQLMFTVGIDFTASNGNPL
TEAVTGQHVMFSVYETHVEW RQIGLPDVANQNTRHPVKFEF VSHEFAINFNPTNPFCSGVDGI QAYSACLPHIRFYGPTNFSPIV HVARFAAQATQQRTATQYFII IITDGVISDMEETRHAVVQASI LPMSIIIVGVGNADFAAMEFLI GDSPHAALPTRGRRQPRDIVQ  21734 52102 A 21862 694 825 QPGLSINIEAGAVPRPFARRYV GNLG*FLFLGVLRCFSSPGSPH  21735 52103 A 21863 1032 1127 FYAKAEAGDRTVA*PHKPKM			1				DPSSLHYINPMGTNEYLSAIWA
RQIGLPDVANQNTRHPVKFEF VSHEFAINFNPTNPFCSGVDGI QAYSACLPHIRFYGPTNFSPIV HVARFAAQATQQRTATQYFII IITDGVISDMEETRHAVVQASI LPMSIIIVGVGNADFAAMEFLI GDSPHAALPTRGRRQPRDIVQ 21734 52102 A 21862 694 825 QPGLSINIEAGAVPRPFARRYV GNLG*FLFLGVLRCFSSPGSPH 21735 52103 A 21863 1032 1127 FYAKAEAGDRTVA*PHKPKM							VGQIIQDYDRPRSFLKPDPWAG
VSHEFAINFNPTNPFCSGVDGI QAYSACLPHIRFYGPTNFSPIV HVARFAAQATQQRTATQYFII IITDGVISDMEETRHAVVQASI LPMSIIIVGVGNADFAAMEFLI GDSPHAALPTRGRRQPRDIVQ 21734 52102 A 21862 694 825 QPGLSINIEAGAVPRPFARRYV GNLG*FLFLGVLRCFSSPGSPH 21735 52103 A 21863 1032 1127 FYAKAEAGDRTVA*PHKPKM							TEA:VTGQHVMFSVYETHVEWG
QAYSACLPHIRFYGPTNFSPIV HVARFAAQATQQRTATQYFII IITDGVISDMEETRHAVVQASI LPMSIIIVGVGNADFAAMEFLI GDSPHAALPTRGRRQPRDIVQ  21734 52102 A 21862 694 825 QPGLSINIEAGAVPRPFARRYV GNLG*FLFLGVLRCFSSPGSPH  21735 52103 A 21863 1032 1127 FYAKAEAGDRTVA*PHKPKM							RQIGLPDVANQNTRHPVKFEFQ
HVARFAAQATQQRTATQYFII IITDGVISDMEETRHAVVQASI LPMSIIIVGVGNADFAAMEFLI GDSPHAALPTRGRRQPRDIVQ  21734 52102 A 21862 694 825 QPGLSINIEAGAVPRPFARRYV GNLG*FLFLGVLRCFSSPGSPH  21735 52103 A 21863 1032 1127 FYAKAEAGDRTVA*PHKPKM							VSHEFAINFNPTNPFCSGVDGIA
IITDGVISDMEETRHAVVQASI LPMSIIIVGVGNADFAAMEFLI GDSPHAALPTRGRRQPRDIVQ 21734 52102 A 21862 694 825 QPGLSINIEAGAVPRPFARRYV GNLG*FLFLGVLRCFSSPGSPH 21735 52103 A 21863 1032 1127 FYAKAEAGDRTVA*PHKPKM							QAYSACLPHIRFYGPTNFSPIVN
LPMSIIIVGVGNADFAAMEFLI GDSPHAALPTRGRRQPRDIVQ 21734 52102 A 21862 694 825 QPGLSINIEAGAVPRPFARRYV GNLG*FLFLGVLRCFSSPGSPH 21735 52103 A 21863 1032 1127 FYAKAEAGDRTVA*PHKPKM							HVARFAAQATQQRTATQYFILL
GDSPHAALPTRGRRQPRDIVQ   21734   52102   A   21862   694   825   QPGLSINIEAGAVPRPFARRYV GNLG*FLFLGVLRCFSSPGSPH   21735   52103   A   21863   1032   1127   FYAKAEAGDRTVA*PHKPKM							IITDGVISDMEETRHAVVQASK
21734         52102         A         21862         694         825         QPGLSINIEAGAVPRPFARRYV GNLG*FLFLGVLRCFSSPGSPH           21735         52103         A         21863         1032         1127         FYAKAEAGDRTVA*PHKPKM							LPMSIIIVGVGNADFAAMEFLD
GNLG*FLFLGVLRCFSSPGSPH 21735 52103 A 21863 1032 1127 FYAKAEAGDRTVA*PHKPKM							GDSPHAALPTRGRRQPRDIVQF
GNLG*FLFLGVLRCFSSPGSPH 21735 52103 A 21863 1032 1127 FYAKAEAGDRTVA*PHKPKM	21734	52102	Α	21862	694	825	QPGLSINIEAGAVPRPFARRYW
							GNLG*FLFLGVLRCFSSPGSPH
	21735	52103	Α	21863	1032	1127	FYAKAEAGDRTVA*PHKPKMF
LSIHHRVAKI							LSIHHRVAKI
21736 52104 B 21864 1 564	21736	52104	В	21864	1	564	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence ( X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21737	52105	Α	21865	43	617	PIIQKVRSHTPKRAPTACTYTVS
						GSFSLPSPGFFSPFPHGTGSLSVS
						QEYLALEEWSPHISARDCTCSG
						LLIELPSTGIFVYGAVTLYRAPF
						QTLPVTHALIQALGCSP\FARRY
					ļ	WGNLG*FLFLGVLRCFSSPGSP
						H*PMDSVNDSVSKHT/EVSPFG
						N\PGYNGSYHLTDAYRRLARPS
		1				SPLT\PGHPPLYG
21738	52106	Α	21866	2969	3593	
21739	52107	Α	21867	1219	1503	HFTDGQNNRQHGCGVRRQAG
						KQVVCKVEHQNQYNHRCVW*
						ESFQVRTHGTGEPVNKTRTSQA
		ĺ				VSHHNQSCQVDQCVPCAVIAG
		l				NVFPRHNVQGQH
21740	52108	Α	21868	193	378	
21741	52109	В	21869	1	1473	
21742	52110	Α	21870	3536	3926	PKMVRWLARKCWNTVSTVRC
						FWMAMPTPVFAPCAAIKTASA
1		ŀ				R*MSWASSR*PNRGCVKSATLR
	1					QFS*VAAMK*PPVAQ*WW\LGR
						NASTAGGDSGAGRSLDDGEPTP
		ŀ				CGMGLEQKRHGNPAGCVAPSR
						WS
21743	52111	Α	21871	140	901	MAMSKVKSITRESWILSTFPEW
1						GSWLNEEIEQEQVAPGTFTMW
						WLGCTGIWLKSEGGTNVCVNL
						WCGTGKQSHGNPLMKQGQQM
						PPMAGVKKLQPNLRTTPFVLDP
						FAIRQIDAVLATHDHNDHIDVN
		1				VAAAVMQNCADDVPFIGPKTC
		1				VDSWIGWGVPKERCIVVKPGD
						VVQVTALEIHAPDAFDRTAPIT
						WLADQKRRRTVKG/MDDAPL*
						QVSYRPSTPPRCCYWSLLSSSV
						YYSKRPFAHGYARLCIPFMD
21744	52112	Α	21872	287	392	
21745	52113	Α	21873	1	492	
21746	52114	A	21874	569	1061	SGTVLDM/SEYSRVQIRKQPYLS
						KGYIPVRQKTKAAWLLKAVTFI
						DLTTLSGDDTSSNIQRLCYKAK
						YPIREDLLKALNMHDKVAAGF
	1					PAGQTHLKTRLEEIRLAVEDGA
]						TEIDVVINRSLVLTGQWEG/LYS
						PVVYSALPCNI*HSRLFLPSGNA
L						LLLWLWCLHS

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NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
}	İ		]	sequence		
21747	52115	A	21875	2	1072	ARSNIRMVTAFRSLPAGSSGAA
		l				RA\MSAHNRGTELDLSWISKIQ
						VNHPAVL\RRAEQIQARRT\VK
		Ì				KDWQAAWLLKAVTFIDLTTLS
!						GDDTSFNI*RLCYKAGYPIREDL
						LKALNMHDKGIT\TAAVCV\YP
1						ARV\CDAVKGTSRPAG/CVNIPV
		1				GISRAAGF\PAG\QTH\LKTRLEE
		İ				I\RLAVEDGSLQKIDV\VINRSLG
		ŀ				A*QAQWE\ALYDEIRQF\RKAC
						GEA\HLKTILA\TGELGTLTNVY
						K\ASMIAMMAGSDF\IKTSTGKE
						TVNATFPVAIVMLRAIRDFFWK
						TGNKIGFKPAGGIRSAKDSLAW
ļ						LSLVKEELGDEWLKP\ELFRIGA
1						ST\LLSDIER\QIYHSCGLGRYGS
						LFMDLSQCL
21748	52116	Α	21876	3	221	
21749	52117	В	21877	49	342	
21750	52118	Α	21878	1	828	
21751	52119	Α	21879	1362	2405	RPPWRHWDGNSSVLVHFQLHF
						LLRPLQTLSLGLEEELLQRGIRA
						RLREHGISLAAYGTIVSAELKG
						GSRPCAVWWPSEPGAGRCPGN
						SFSCGNSQCVTKVNPECDDQED
		ŀ				CSDGSDEAHCECGLQPAWRMA
						GRIVGGMEASPGEFPWQASLRE
						NKEHFCGAAIIN/DQPLRS*NHR
						FQDPTKWVAYVVRPTSAARRP
		1				APCGPSKKA/WISGWGYLK/EG
1			:			LP*ASPRPAMVKPEVLQKA/DW
						ELLDQALCASLYGHSL/TDRMV
						CAGYLDGKVDSCSVLFLFRGTI
						SFFFSEDRNLIRYVLLYSIFVVN
						CISGKPQPGPPVLRALCQHPHQI
						DAGPQYRASCWVTVPKLQGIF
21752	52120	Α	21880	600	904	SVSLLDEETALPFSWDELRLSA
						LLDT*AAFQEIELAFQDPQCVG
						LLTAP/YADAAFSGSTDIAMFLL
						VPILSARKPASPRSS*TFSHLATE
						PLPASIK*AAA
21753	52121		21881	393	572	
21754	52122	A	21882	165	432	PGMAVSYEKISKYPVEFLPAQA
						NIAGHNDAPEEREPFQW*LWLL
						LPNF*LQEKCSPTLCHFLSRRTQ
				_		SEFPSSTAAGNLSVSLLYPCKS
21755	52123	В	21883	1	1011	

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1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21756	52124	A	21884	783	1234	YYRRCGAASPHGRDPVQIG*TS
				<b>'</b>		SQPR*SAYRN/PASVIPVARHSA
						RAPAILRP*VVVAERSERSIPGL
						RATPAEIRTTSAPSSAGVASSPE
1						KPLIFTAIDSHTGSHWSDIIHVC
						AKKLQNYVPKLNASLLLTPSIY
						VTRTTRSGPIPPARQ
21757	52125	A	21885	249	398	GYFDIFSYDTAIPGQSAEQDAR
						ED*AF*MPDRAYSQPCSSLVFP
						KGCKK
21758	52126	Α	21886	309	845	CGAREERGRGCHHVHSQLRGP
						GPAPGQVQQPQLVPAPVVPELP
į						GLRGLSWQARRTRARHRQGQV
						QDHQPDPTVHAQLRGVQLGEA
						PLQLHHGD*DHRAP*GYFDIFS
						YDTAIPGQSAEQDARED*AF*M
						PDRAYSQPCSSLVFPKGCPVPG
		İ				RGCAAGVQAAGAAHPPLDHPA
						LQPLQGR
21759	52127	A	21887	1236	2100	DLLCGRGCQGRRPRHLQPQTEL
21739	32127	^	21007	1230	2100	EAPEVPAERPGRGPAVAARSDP
						RAQAEDVPGYNPLSGKPAKQQ
						VHEGHHPGAPGHRRPGGAAVP
İ						AGGPHRCLPAQPARPCGGAAEP
						RGGPAGAH*ARGSRAAGEGQP
ŀ						WWPSPRLGAGPPAAPVHAAPG
						SHLLPGHHLPTQPSALAAQ\PL
						AEEPREQ/PGAGGPQPGSPTLRG
			•			VEVVPLPHAAGGAGAVPVGGA
						ACPTAPHPAASAPAPVLLSQLG
		1				TQHPPRRDPPHCSCAGIQDSGW
						AGPPALWSLLHVAKPAQARRP
						CALLHPGV
21760	52128	Α	21888	1931	2185	AKAFSFFRYISRVSRNRTCLSRS
						SMRRSFDSSNADAAFSGSTDIA
						MFLLVPILSARKPASPRSS*TFS
						HLATEPLPASIK*AAA
21761	52129	Α	21889	1254	1760	
21762	52130	Α	21890	1	2001	SIRRPKRARPIRIKKIPEQQG/PQ
						PAVLPVQTVATRDRELPQMP/P
						VGPETLKRDPPVRAMTIPATAA
		l				VYKPYWGATPLPIA\KPSPAG*Q
		<u> </u>				*SRRFSKGYFGFD
21763	52131	Α	21891	195	545	HSPGSFYSRRDAQSPAPAIRAA
		1				V*AEVQ*TPLCHDRITHFPEDW
		i		1		RGWQRS*F*PAACRVAGSLPVP
						VSPQIQHPYCRSSAQYLPCGAG
						LPLPGFDRLCPRSLFQGLAVTG
		1		1		GSCGPS
L	l		l	<del></del>		

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ļ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21764	52132	Α	21892	825	1299	YSCRTEHPPRLATAEYRHHQRA
						ADAGR*RHAGVAFYPIKERYR
						QPEPAFSYQ/PRRWSASRNRRPS
						R*RMRRFPANSSTPFPPAPPLRR
						KRVASLIGGRMLRLTGAGIAEE
						RMIAPQLPECILHELTERPHPFP
						LGIDLILTCGERLLAIPRTTHVE
21765	52133	Α	21893	1460	1810	RATHVLTWVRGTPKRAEMILG
						SSRSMSAAAVTVLISPATAQRW
						ITRAP*PSPIRSPSVSRLTATRPSS
						ITKARALSPMAAPALRSTVTTP
						PRITVEKPLLMEKIPRVPKSRAI
						SAL
21766	52134	A	21894	231	704	QSQKPDTRGRLFVAKAPPCVQR
						HKVRWLRLVMITSAHYLPLRR*
						*PAHPASGGCCPNRAKYTQRPA
						E*ISYRPD*EAVR*SRAPAKVTQ
						SNPASRVALRRPAPRAVHQATII
						PVDESTLLPAPLAVSARQKVRQ
		l				AIPLRDSPVRRPTLDVRHRATG
			i.			L
21767	52135	A	21895	1291	1925	RQLLRWSNR*IAR*SLTQPARRF
						PAYHYAHPAAAAAPGDPLGAD
						LQLCARHYHRQRHCVPRHAGY
						PRSDGLHPQPRGRWRIRCSDCL
						RVPF
21768	52136	Α	21896	1	445	MFISGLDDFFIDVVYWVRRIKR
						KLSVYRRYPRMSYRELYKPDE
						KPL/DDYGSGVE*NGRHRQYGR
						AGGDHARLRKLSYLCWHLPQR
						PRYSA*C*RSVRSLPECA*GSLR
ŀ						ASWP/HPAKPTV*TTCWTPSPN
						LSVAPISLLLVLFCMTPKM
21769	52137	Α	21897	79	377	SRKSGISIGAVKRPTH*GS*KAR
						SIC*NAAYVSKKAESLSSFPEKS
						ESTQRVKAVLSS**ASQGCGDT
						AQYVLLSFKGSQQA*QVCGR*
						KAY**DLSRE
21770	52138	Α	21898	1282	1535	GSTYEIASATPSPSAKPSNRSPSP
		1				MSRPQSNPKPNSLHSC*P*GNL
						RSTPESGK*ATPPPPPAYTLVLF
						KLANPQLPEESPWAF
21771	52139	Α	21899	3775	4031	CRRTAASRTPRKRTQTAPTHAT
						NKPQKPQTSTDRGNRREPEKTE
		1				PS*HLRPNPQEPGRRGQTQQ\EP
						APRHQPPKPQGEPTTPQKQ
L				<del></del>	L	

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NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21772	52140	Α	21900	1	2595	MESHTVTDYNFGRLLNESFAN
21772	32140	(`	21700	1	2373	CETDRNYLGIFKCGFWLSGTRK
						SAFLTKSQVTLTRPVDHTLRNE
						VLGSTKYQVPGYRRKTNTNLE
						ESNVLIGYDVAPGGGKQIDGKK
						AEGRELMVPVAGRPLCRSSSM
						ELLLLPLHIPREQMLLADAYW
						NLHQSCQSKTSPDTAKCPLDRC
						PRVFKSQSYKASTFDGLSMDTE
						VIKSNGKQGWRQRRLANMKDI
						DKQTQAATFVLSQGMQRSGND
						RIILVTGASDGIGREAAMTYAR
						YGATVILLGRNEEKLRQVASHI
						NEETGRQPQWFILDLLTCTSEN
						CQQLAQRIAVNYPRLDGVLHN
			!			AGLLGDVCPMSEQNPQVWQD
				ļ		VMQLFRAGIINADPGAFFVTGF
						RRYCCFTLCCRSFRFGFYRRRF
						l I
					]	RHRSCRQRAMRHSGTGQGQGN
						EQCFNSVFHTVFSLFVNSGITLT
						AVKTVNKWQCSEMEKAAKAT
						FSCYSCNAFFCEAASGLLLFQRI
						ANFDFRLFTQVTHQSFKDDLGI
						KMETTKPSFQDVLEFVRLFRRK
						NKLQREIQDVEKKIRDNQKRVL
						LLDNLSDYIKPGMSVEAIQGIIA
						SMKGDYEDRVDDYIIKNAELSK
j				ļ		ERRDISKKLKAMGEMKNGEAN
						RKERSSMFRDRPLKRMKSKRD
						DDSYDEDVEDDEGVGEVRVHR
		ĺ				VNHAPANAQEHEAARPSPQHQ
						YQPPYASAQPRQPVQQPPEAQV
21773	52141	Α	21901	1	96	
21774	52142	Α	21902	105	430	ASIFVMLVRGAEPMEKRQQRG
						LFTVPGLLLAFCSHVLSCVNPLI
						PVDNPYYPPLSELIPAPPQPNDR
						AQRVSERGSGRAPNTQTASPRA
L						LADSLMQLAR\QVSRLESGQ